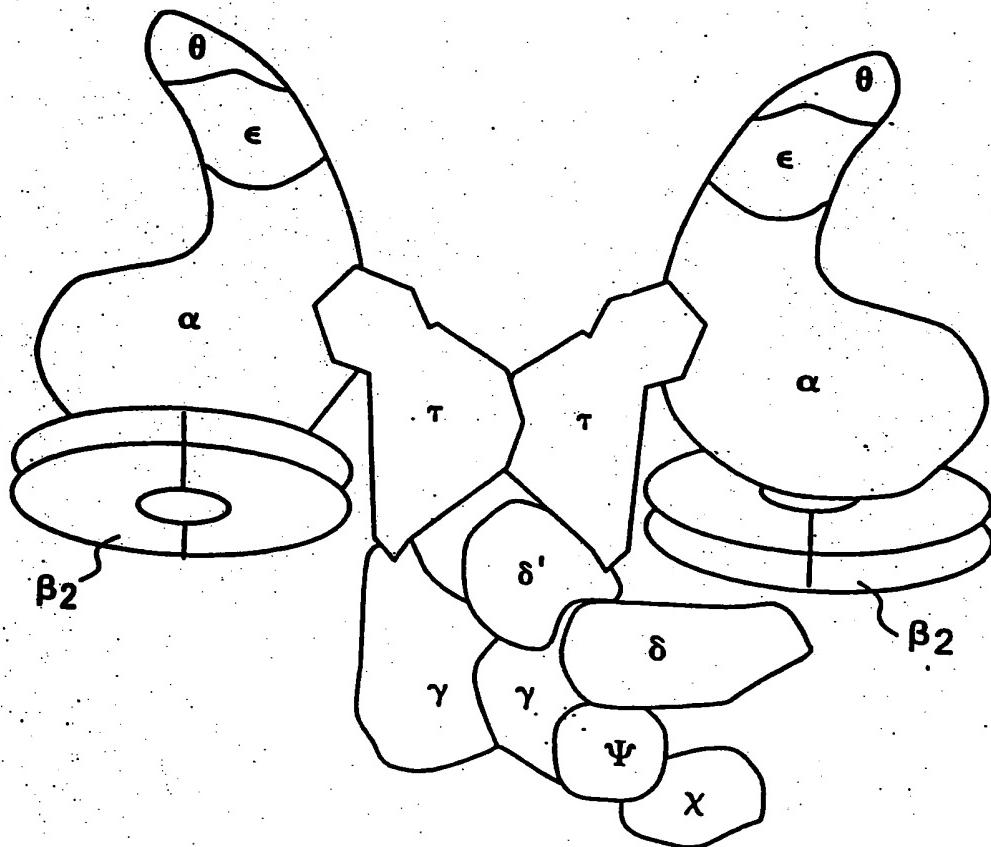


FIG.1



ATP binding

2

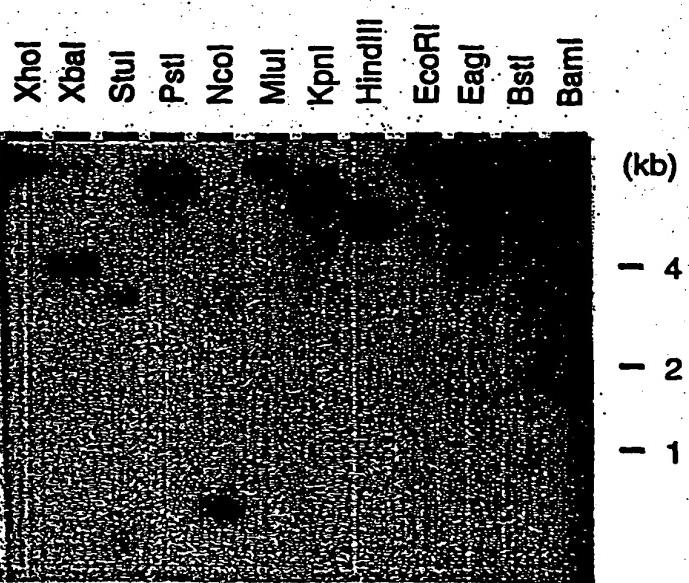


FIG.3

FIG. 4A-1

TGS CTS CTC CTC GGS CTC GTG
 ACC CTG GAG CCC CCG CCC CAC GTC CTC GTC GTC ACC ACC GAG CCC GAG ACG 600
 thr leu glu pro pro his val leu phe val phe ala thr thr glu pro glu arg (157)

 ATG CCC CCC ACC ATC CTC TCC CGC ACC CAG CAC TTC CGC TTC CGC CTC ACG GAG GAG 660
 met pro pro thr ile leu ser arg thr gln his phe arg phe arg arg leu thr glu glu (177)

 GAG ATC GCC TTT AAG CTC CGG CGC ATC CTG GAG GCC GTG GGG CGG GAG GAG GAG 720
 glu ile ala phe lys leu arg arg ile leu glu ala val gly arg glu ala glu glu glu (197)

 GCC CTC CTC CTC GCC CGC CTG GCG GAC CGG CCC CTC AGG GAC GCG GAA AGC CTC CTC CTC CTC CTC 780
 ala leu leu leu ala arg leu ala asp gly ala leu arg asp ala glu ser leu leu (217)

 GAG CGC TTC CTC CTC CTG GAA GGC CCC CTC ACC CGG AAG GAG GTG GAG CGC CGC CTA GGC 840
 glu arg phe leu leu glu gly pro leu thr arg lys glu val glu arg ala leu gly (237)

 TCC CCC CCA GGG ACC GGG GTG GCC GAG ATC GCC TCC CTC GCG AGG GGG AAA ACG GCG 900
 ser pro gly thr gly val ala glu ile ala ser leu ala arg gly lys thr ala (257)

 GAG GCC CTG GGC CTC CGG CGC CTC TAC GGG GAA TAC GCC CGC CCG AGG AGC CTC GTC GTC GTC 960
 glu ala leu gly leu ala arg arg leu tyr gly glu gly tyr ala pro arg ser leu val (277)

 TCG GGC CTT TTG GAG GTG TTG CGG CTC TAC GCC GCC TTG GGC CTC CGC GCA ACC ACC 1020
 ser gly leu leu glu val phe arg glu gly leu tyr ala ala phe gly leu ala gly thr (297)

 CCC CTT CCC CCG CCC CAG GCC CTG ATC GCC ACC GAC GAG GCC ATG 1080
 pro leu pro pro gln ala leu ile ala met thr ala ala asp glu ala met (317)

FIG. 4A-2

GAG CGC CTC GCC CGC TCC GAC GCC TTA AGC CTG GAG GTG GCC CTC CTG GAG GCG GGA 1140
glu arg leu ala arg ser asp ala leu ser leu glu val ala leu leu glu ala gly (337)

AGG GCC CTG GCC GGC GAG GCG CTA CCC CAG CCC ACG GGC GCT CCT TCC CCA GAG GTC GGC 1200
arg ala leu ala ala glu ala leu pro gln pro thr gly ala pro ser pro glu val gly (357)

CCC AAG CCG GAA AGC CCC CCG ACC CCC CCA AGG CCC GAG GAG GCG CCC GAC CTG CTG 1260
pro lys pro glu ser pro pro pro pro arg pro pro glu glu ala pro asp leu (377)

CGG GAG CGG TGG CGG GCC TTC CTC GAG GCC CTC AGG CCC ACC CTA CGG GCC TTC GTG CGG 1320
arg glu arg trp arg ala phe leu glu ala leu arg pro thr leu arg ala phe val arg (397)

GAG CGC CCG GAG GTC CGG GAA GGC CAG CTC TGC GCT TTC CCC GAG GAC AAG GCC 1380
glu ala arg pro glu val arg glu gly gln leu cys leu ala phe pro glu asp lys ala (417)

TTC CAC TAC CGC AAG GGC TCG GAA CAG AAG GTG AGG CTC CTC CCC CTG GCC CAG GCC CAT 1440
phe his tyr arg lys ala ser glu gln lys val arg leu leu pro leu ala gln ala his (437)

frameshift site

TTC GGG GTG GAG GTC GTC CTC GTC CTG GAG GGA GAA AAA AGC CTG AGC CCA AGG 1500
phe gly val glu glu val val leu leu glu gly glu lys lys ser leu ser pro arg (457)

FIG. 4B - 1

CCC CGC CCG CCC CCA CCT CCT GAA GCG CCC GCA CCC CCC GGC CCT CCC GAG GAC GAG GTC	1560					
pro arg pro ala pro pro glu ala pro pro glu pro pro glu glu val	(477)					
GAG GCG GAG GAA GCG GCG GAG GAG GCG CCC GAG GAG GCG TTT AGG CGG GTG GTC CGC CTC	1620					
glu ala glu ala ala glu glu ala pro glu glu ala leu arg arg val val arg leu	(497)					
CTG GGG CGG CGG GTG CTC TGG GTG CGG CGG CCC AGG ACC CGG GAG GCG CCG GAG GAA	1680					
leu gly gly arg val leu trp val arg arg pro arg thr arg glu ala pro glu glu glu	(517)					
CCC CTG AGC CAA GAC GAG ATA GGG GGT ACT GGT ATA TAA	TGGGGCCATG	ACGGGGACAC	1740			
pro leu ser gln asp glu ile gly gly thr gly ile *			(529)			
CGACCTCGGA CAAGAGACCG	TGGACAAACAT	CCTCAAGGCC	CTCCGGCGTA	TTGAGGGCCA	1820	
GGTGGGGGG	CTCCAGAAGA	TGGTGCCCCGA	GGGGCGCCCC	TGGGACGGGG	TCCTCACCCA	1880
GATGACCGCC	ACCAAGAAGG	CCATGGAGGC	GGGGCCACC	CTGATCCTCC	ACGAGTTCT	1940
GAACGTTCTGC	GCCGCCGAGG	TCTCCGAGGG	CAAGGTGAAAC	CCCAAGAAC	CCGAGGAGAT	2000
CGCCACCATG	CTGAAGAACT	TCATCTA				

FIG.4B-2

FIG. 4C

51

GTG	AGC	GCC	TTC	TAC	CGC	CGC	TTC	CCC	CTC	ACC	TTC	CAG	TTC	GAG	GTG	GTG
GGG	CAG	GAG	CAC	GTG	AAG	GAG	CCC	CTC	CTC	AAG	GGG	CGG	GGG	AGG	CTC	GCC
GCC	TAC	CTC	TTC	GGG	CCC	AGG	GGG	GGG	GGG	AAG	ACC	ACC	ACG	GGG	CTC	CTC
ATG	GGC	GTG	GGG	TGC	CAG	GAA	GAC	CCC	CCT	TGC	GGG	GTC	TGC	CCC	CAC	TGC
GtG	CAG	AGG	GGC	GCC	CAC	CCG	GAC	GTG	GTG	GAC	ATT	GAC	GCC	AGC	AAC	TCC
GAG	GAC	GTG	GGG	GAG	CTG	AGG	GAA	AGG	ATC	CAC	CTC	CCC	CTC	TCT	CCC	AGG
GTC	TTC	ATC	CTG	GAC	GGC	CAC	ATG	CTC	TCC	AAA	AGC	GCC	TTC	AAC	GCC	CTC
ACC	CTG	GAG	GCC	CCC	CCG	CAC	GTC	CTC	TTC	GTC	TTC	GCC	ACC	ACC	GAG	AGG
ATG	CCC	ACC	ATC	CTC	TCC	CGC	ACC	CAG	CAC	TTC	CGC	TTC	CGC	CTC	ACG	GAG
GAG	ATC	GCC	TTT	AAG	CTC	CGG	CGC	ATC	CTG	GAG	GCC	GTG	GGG	GGG	GAG	GAG
GCC	CTC	CTC	CTC	CTC	CTC	CGC	CGC	CTG	GAC	GGG	GCC	CTT	AGG	GAC	GGG	GAG
GAG	CGC	TTC	CTC	CTC	CTC	CTG	GAA	GGC	CCC	CTC	ACC	CGG	AAG	GAG	GGG	GAG
TCC	CCC	CCA	GGG	ACC	GGG	GTG	GCC	GAG	ATC	GCC	GCC	TCC	CTC	GGG	AAA	AGC
GAG	GCC	CTG	GGG	CTC	GGC	CGG	CGG	CTC	TAC	GGG	GAA	GGG	TAC	GCC	CCG	AGG
TCG	GGC	CTT	TTG	GAG	GTG	TTG	GAA	GGC	CTC	TAC	GCC	TCC	GGC	TTC	GGG	AGC
CCC	CTT	CCC	GCC	CCC	CCC	CAG	GCC	CTG	ATC	GCC	GCC	ATG	ACC	GGC	CTG	GCC
GAG	CGC	CTC	GCC	CGC	TCC	GAC	GCC	TTA	AGC	CTG	GAG	GTG	GCC	CTC	CTG	GAG
AGG	GCC	CTG	GCC	GCC	GCC	GAG	CTA	CCC	CCC	ACG	GGC	GCT	CCT	TCC	CCA	GAG
CCC	AAG	CCG	GAA	AGC	CCC	CCG	ACC	CGG	ACC	CGG	CCC	CCA	AGG	CCC	CTG	GCC
CGG	GAG	CGG	TGG	CGG	GCC	TTC	CTC	GAG	GCC	CTC	AGG	CCC	ACC	CTA	CGG	CCC
GAG	GCC	CGC	CGG	GAG	GTG	GTC	GAA	GAG	GTC	GTC	GTC	GAG	AAA	AAA	AGC	TTC
TTC	CAC	TAC	CGC	AAG	GCC	TCG	GAA	CAG	AAG	GTG	AGG	CTC	CTC	CCC	CTG	GCC
TTC	GGG	GTG	GAG	GAG	GTC	GTC	GTC	CTG	GAG	GGA	GAA	AAA	AAA	AGC	CTG	AGG
CCC	CGC	CCG	GCC	CCA	CCT	GAA	GCG	CCC	GCA	CCC	CGG	CCT	CCC	GAG	GAG	GAG
GAG	GCG	GAG	GAA	GCG	GCG	GAG	GAG	CCC	GAG	GCC	TTC	AGG	CGG	GTC	CGC	CTC
CTG	GGG	GGG	GGG	CTC	TGG	GTG	CGG	CGG	CCC	AGG	ACC	CGG	GAG	CCG	GAG	GAA
CCC	CTG	AGC	CAA	GAC	GAG	ATA	GGG	GGT	ACT	GGT	ATA	TAA	(1590)			

FIG. 4D

Met ser ala leu tyr arg arg phe arg pro leu thr phe gln glu val val gly gln glu 20
his val lys glu pro ile arg glu gly lys thr thr ala arg leu leu ala gln ala tyr leu 40
phe ser gly pro arg gly val gly val cys pro his cys gln ala val met ala val 60
gly cys gln gly glu asp pro pro cys gly val asp ile asp ala ala ser asn ser val glu arg 80
gly ala his pro asp val val ser ala pro leu ser ala pro arg lys val phe ile 100
arg glu leu arg glu arg his leu ser lys ser ala phe asn ala leu leu lys thr leu glu 120
leu asp glu ala his met leu ser lys ser ala phe arg arg leu thr glu glu glu ile ala 140
glu pro pro his val leu phe val phe ala thr thr glu pro glu arg met pro pro 160
thr ile leu ser arg thr gln his met leu ser lys ser ala phe arg arg leu thr glu glu glu ile ala 180
phe lys leu arg arg ile leu glu ala val gly arg glu ala glu glu glu ala leu leu 200
leu leu ala arg leu ala asp gly ala leu arg asp ala glu ser leu leu glu arg phe 220
leu leu leu glu gly pro leu thr arg lys glu val glu arg ala leu gly ser pro pro 240
gly thr gly val ala glu ile ala ala ser leu ala arg gly lys thr ala glu ala leu 260
gly leu ala arg arg leu tyr gly glu gly tyr ala pro arg ser leu val ser gly leu 280
leu glu val phe arg glu gly leu tyr ala phe gly leu ala gly thr pro leu pro 300
ala pro pro gln ala leu ile ala ala met thr ala leu asp glu ala met glu arg ala leu 320
ala arg arg ser asp ala leu ser leu glu val ala leu glu ala gly arg ala leu 340
ala ala glu ala leu pro gln pro thr gly ala pro ser pro glu val gly pro lys pro 360
glu ser pro pro thr pro glu pro pro arg pro glu glu ala pro asp leu arg glu arg 380
trp arg ala phe leu glu ala leu arg pro thr leu arg ala phe val arg glu ala arg 400
pro glu val arg glu gly gln leu cys leu ala phe pro glu asp lys ala phe his tyr 420
arg lys ala ser gln gln lys val arg leu pro leu ala gln ala his phe gly val 440
glu glu val val leu val glu gly glu lys ser leu ser pro arg pro arg pro 460
ala pro pro glu ala pro pro glu glu glu val val arg val val arg leu leu gly gly 480
glu ala ala glu glu ala pro glu glu glu ala leu arg arg val val arg leu leu gly gly 500
arg val leu trp val arg pro arg thr arg glu ala pro glu glu glu pro leu ser 520
gln asp glu ile gly gly thr gly ile 529

FIG. 4E

Met ser ala leu tyr arg arg phe arg pro leu thr phe gln glu val val gly gln glu 20
his val lys glu pro leu leu lys ala ile arg glu gly arg leu ala gln ala tyr leu 40
phe ser gly pro arg gly val gly lys thr thr ala arg leu leu ala met ala val 60
gly cys gln gly glu asp pro pro cys gly val cys pro his cys gln ala val gln arg 80
gly ala his pro asp val asp ile asp ala ala ser asn ser val glu asp val 100
arg glu leu arg glu arg ile his leu ala pro leu ser ala pro arg lys val phe ile 120
leu asp glu ala his met leu ser lys ser ala phe asn ala leu leu lys thr leu glu 140
glu pro pro his val leu phe val phe ala thr thr glu pro glu arg met pro pro 160
thr ile leu ser arg thr gln his phe arg phe arg glu arg leu thr glu glu ile ala 180
phe lys leu arg arg ile leu glu ala val gly arg glu ala glu glu glu ala leu leu 200
leu leu ala arg leu ala asp gly ala leu arg asp ala glu ser leu leu glu arg phe 220
leu leu glu gly pro leu thr arg lys glu val glu arg ala leu gly ser pro pro 240
gly thr gly val ala glu ile ala ala ser leu ala arg gly lys thr ala glu ala leu 260
gly leu ala arg arg leu tyr gly glu gly tyr ala pro arg ser leu val ser gly leu 280
leu glu val phe arg glu gly leu tyr ala ala phe gly leu ala gly thr pro leu pro 300
ala pro pro gln ala leu ile ala met thr ala ala met glu arg leu 320
ala arg arg ser asp ala leu ser leu glu val ala leu glu ala gly arg ala leu 340
ala ala glu ala leu pro gln pro thr gly ala pro ser pro glu val gly pro lys pro 360
glu ser pro pro thr pro glu pro pro arg pro glu glu ala pro asp leu arg glu arg 380
trp arg ala phe leu glu ala leu arg pro thr leu arg ala phe val arg glu ala arg 400
pro glu val arg glu gly gln leu cys leu ala phe pro glu asp lys ala phe his tyr 420
arg lys ala ser glu gln lys val arg leu leu pro leu gly glu lys lys pro asp pro 440
glu glu val val leu val leu val leu gly glu lys lys pro asp pro lys ala pro pro 460
gly pro thr ser 464

FIG.4F

Met ser ala leu tyr arg arg phe arg pro leu thr phe gln glu val val gly gln glu 20
his val lys glu pro leu leu lys ala ile arg glu gly arg leu ala gln ala tyr leu 40
phe ser gly pro arg gly val gly lys thr thr thr ala arg leu leu ala met ala val 60
gly cys gln gly glu asp pro pro cys gly val cys pro his cys gln ala val gln arg 80
gly ala his pro asp val val asp ile asp ala ala ser asn ser val glu asp val 100
arg glu leu arg glu arg ile his leu ser lys ser ala phe asn ala leu leu lys val phe ile 120
leu asp glu ala his met leu ser arg thr glu pro glu arg met pro pro 140
glu pro pro his val leu phe val phe ala thr thr glu pro glu arg 160
thr ile leu ser arg thr gln his phe arg phe arg arg leu thr glu glu ile ala 180
phe lys leu arg arg ile leu glu ala val gly arg glu ala glu glu ala leu leu 200
leu leu ala arg leu ala asp gly ala leu arg asp ala glu ser leu leu glu arg phe 220
leu leu glu gly pro leu thr arg lys glu val glu arg ala leu gly ser pro pro 240
gly thr gly val ala glu ile ala ala ser leu ala arg gly lys thr ala glu ala leu 260
gly leu ala arg arg leu tyr gly glu gly tyr ala pro arg ser leu val ser gly leu 280
leu glu val phe arg glu gly leu tyr ala ala phe gly leu ala gly thr pro leu pro 300
ala pro pro gln ala leu ile ala met thr ala leu asp glu ala met glu arg leu 320
ala arg arg ser asp ala leu ser leu glu val ala leu glu ala gly arg ala leu 340
ala ala glu ala leu pro gln pro thr gly ala pro ser pro glu val gly pro lys pro 360
glu ser pro pro thr pro glu pro pro arg pro glu ala pro asp leu arg glu arg 380
trp arg ala phe leu glu ala leu arg pro thr leu arg ala phe val arg glu ala arg 400
pro glu val arg glu gly gln leu cys leu cys lys val arg leu leu pro leu his phe his 420
arg lys ala ser glu gln lys lys lys lys lys lys ala his phe gln glu lys 440
glu glu val val leu val leu glu gly glu lys lys ala 454

E. coli	ATP site	MSYQVLARKWKRPTFADVGQEHVLTALANGLSLGRIHHAYLFSGTRGVGKTSIARLLAK	60
H. inf.	K.	II.....KDN.L...	60
B. sub.	A.Y.VF..R.E...	IITKT.Q.A.LQQKFS..P.T...A.KIF...	60
C. cres.	DA.T...Y.R..E.LI...	AMVRT..AF.T..A.FMLT.V...TT...R	113
M. gen.	-MH..FYQ.Y..IN.KQTL...	SIRKI.W.AINRDKLPGNC.I...E..T...TF.KII...	59
T. th.	--VSA.Y.RF..L.QE...	KEP.LKAIRE..LAQ..P..TT..M	58

E. coli	RGRFKVYLIDEVHMLSRHSFNALLKTLLEEPPEHKFELLATTDPQKLPVTLISRCLQFHLLK	176
H. inf.	V.....Y.....	176
B. sub.	AVTY...I.....IGA.....	176
C. cres.	EA.Y...I.....TAA.....	233
M. gen.	TEKK...IL..A...TTQ.WGG.....	175
T. th.	SAPP...FIL..A...KSA.....	172

FIG. 5A

FIG. 5B

E. coli	ALDVEQIRHQLEHILNEEHIAHEPRALQLLARAAEGSLLRDLSSLTQATASGDGQ--VST	234
H.inf.	...ET..SQH.A...TQ.N.PF.DP..VK..K..Q..I..S.....M..R.--TN	234
B.sub.	RITSQA.VGRMNK.VDA.QLQV.EGS.EII.S.H.GM....SFSGDT--LKV	234
C.cres.	RVEPDVLVKHFDR.SAK.GARI.MD..A.I.....V..G...L...VQTERGQT.TS	293
M.gen.	KITSDL.LER.ND.AKK.K.KI.KD..IKI.DLSQ.....G..L..LAI.LIVKKL.LL	235
T.th.	R.TE.E.AFK.RR..EAVGREA.EE..L...L.D.A...E..LERFLLLEGPP--LTR	229
E. coli	QAVSAMIGTLDDQALSILVEAMVEANGERVMALINEAAARGIEWEALLIVEMGLLHRIAM	294
H.inf.	NV..N..L...NYSDILY.LHQG..LL.RTLQRV.DAAGD.DK..G.CAEK..Q..L	294
B.sub.	EDALLIT.AVSQLYIGK.AKSLHDK.VSDALETL..LLQQ.KDPAK.IED.IFYFRDMLL	294
C.cres.	TV.RD..LA.RS.TIA.Y.HVMAGKTCKDALEGFRALWGF.ADPAVVMLDV.DHC.AS.V	353
M.gen.	MLKKHLISLLEMQNL.L.KOFYQ.I	260
T.th.	KE.ERA..SPPGTGVAEIAASLARGKTAEAALG.ARRLYGE.YAPRS.VSGI.EVFREGY	289

FIG. 6

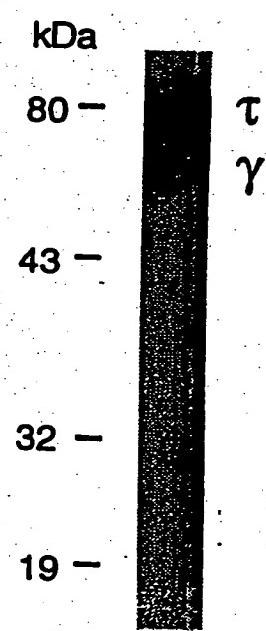
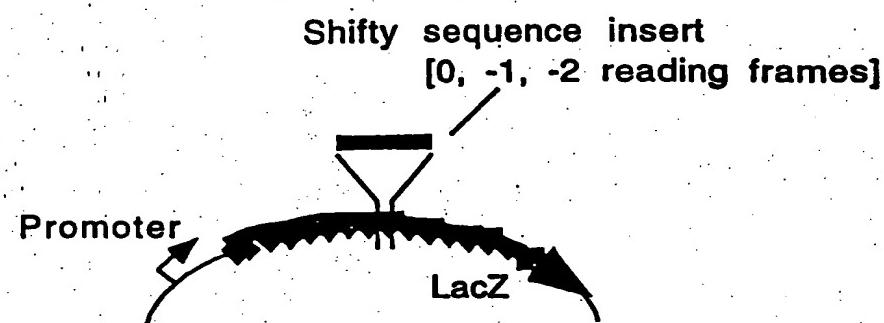


FIG. 7

FIG.8A



	Reading frame	Blue	White
Shifty sequence	0	+	
	- 1	+	
	- 2	+	
Mutant sequence	0	++	
	- 1		+
	- 2		+

FIG.8B

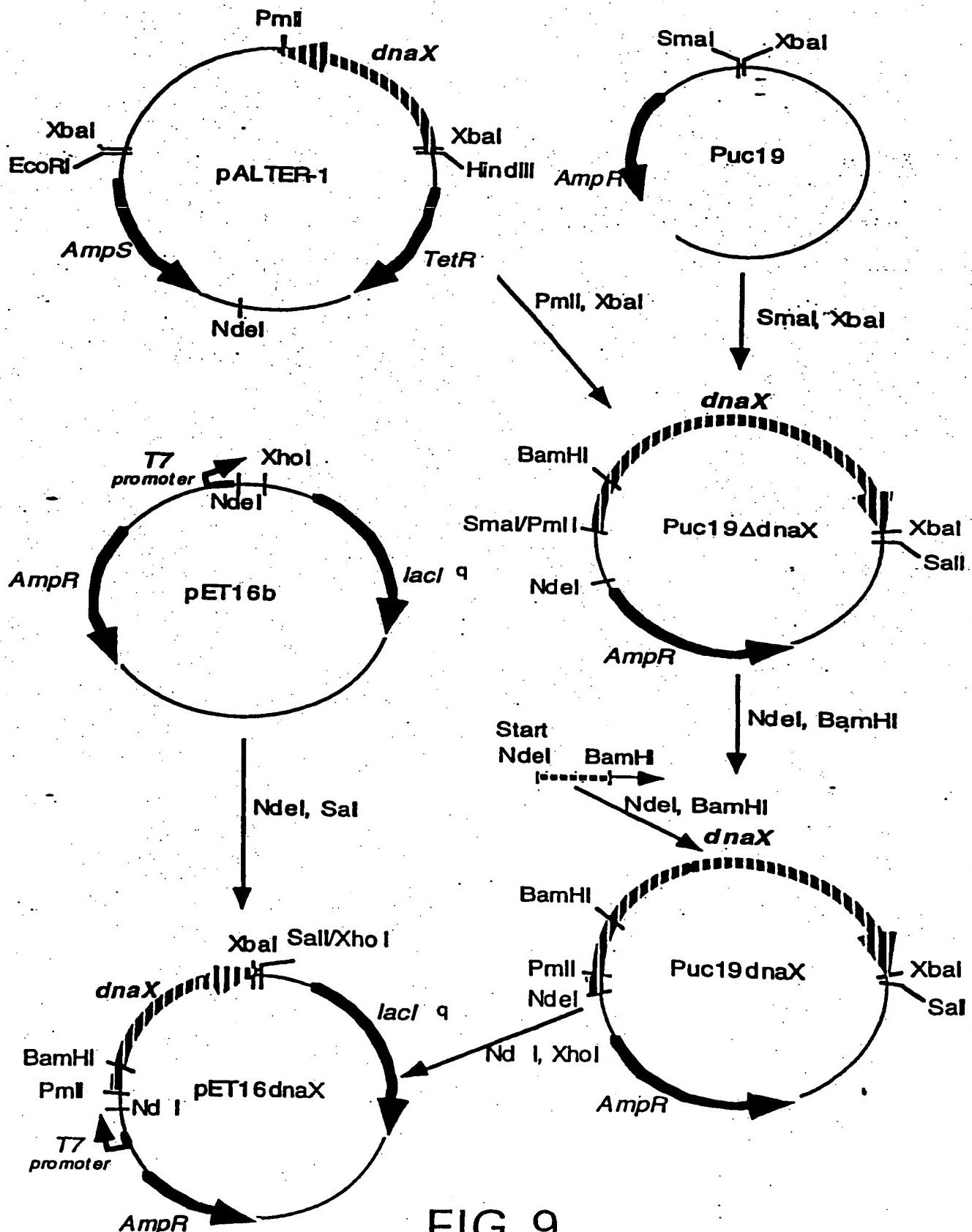


FIG. 9

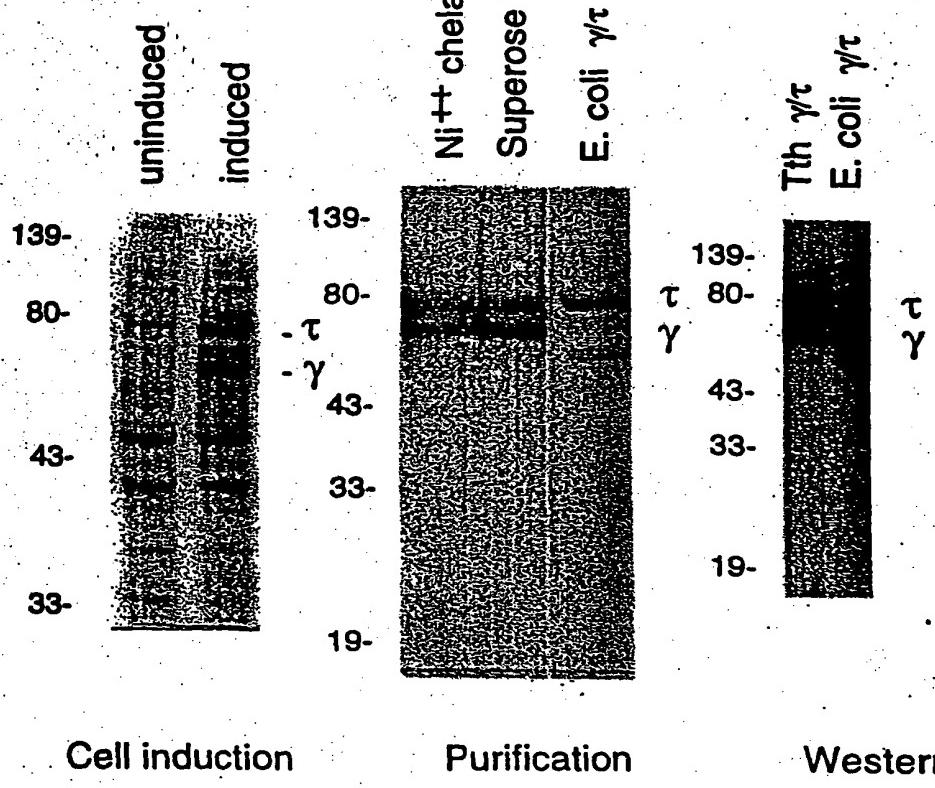


FIG.10A FIG.10B FIG.10C

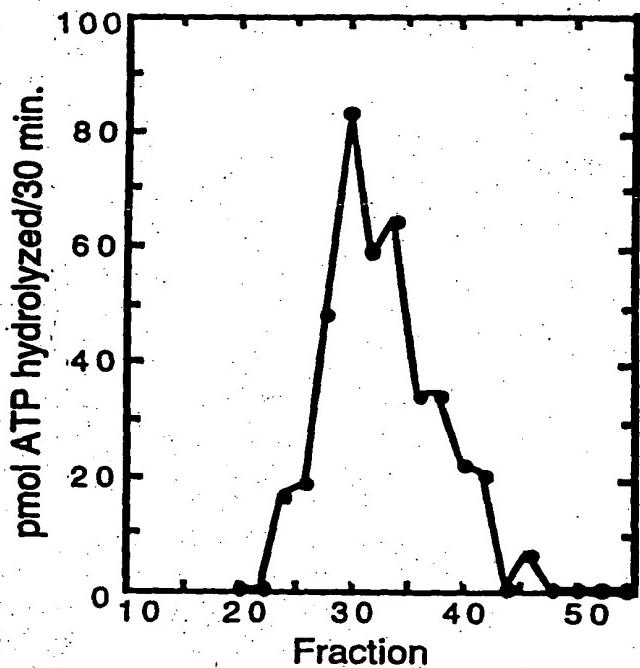


FIG. 11A

kDa 670
↓
Frxn: 18 20 22 24 26 28 30 32 34 36 38 40 42 44 46 48 50 52 54 56



FIG. 11B

FIG. 12A

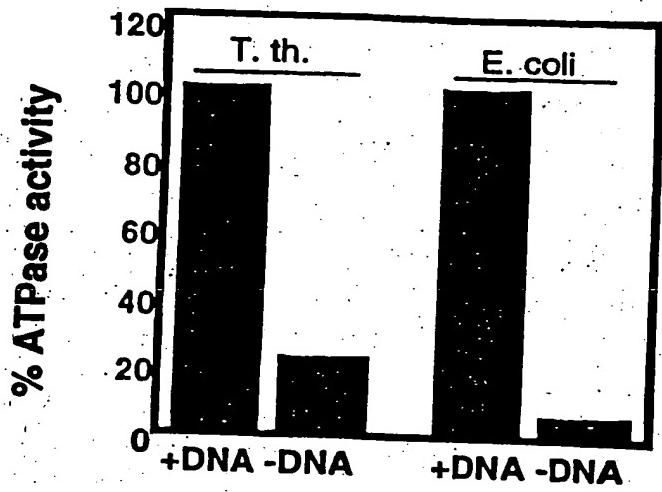


FIG. 12B

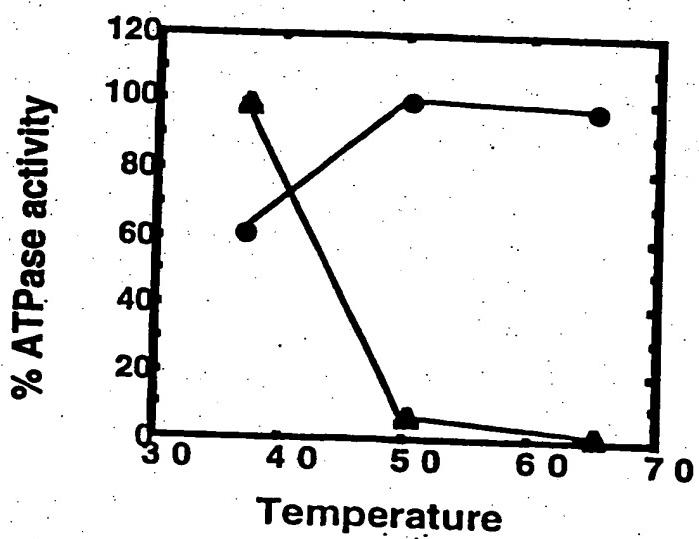


FIG. 12C

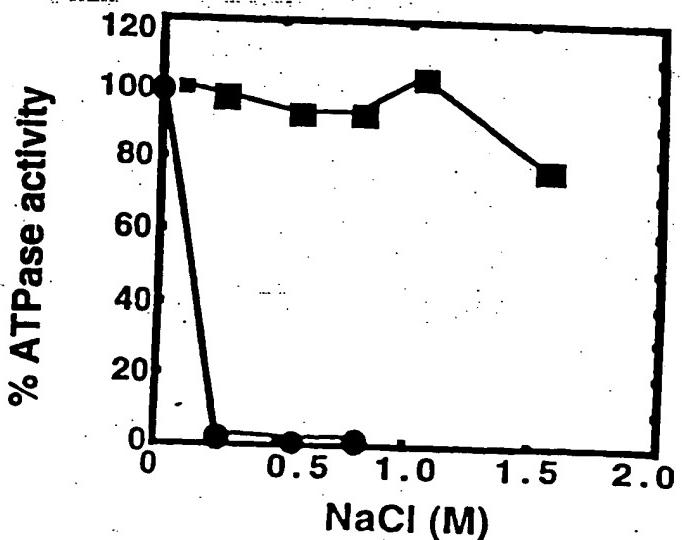


FIG.13A

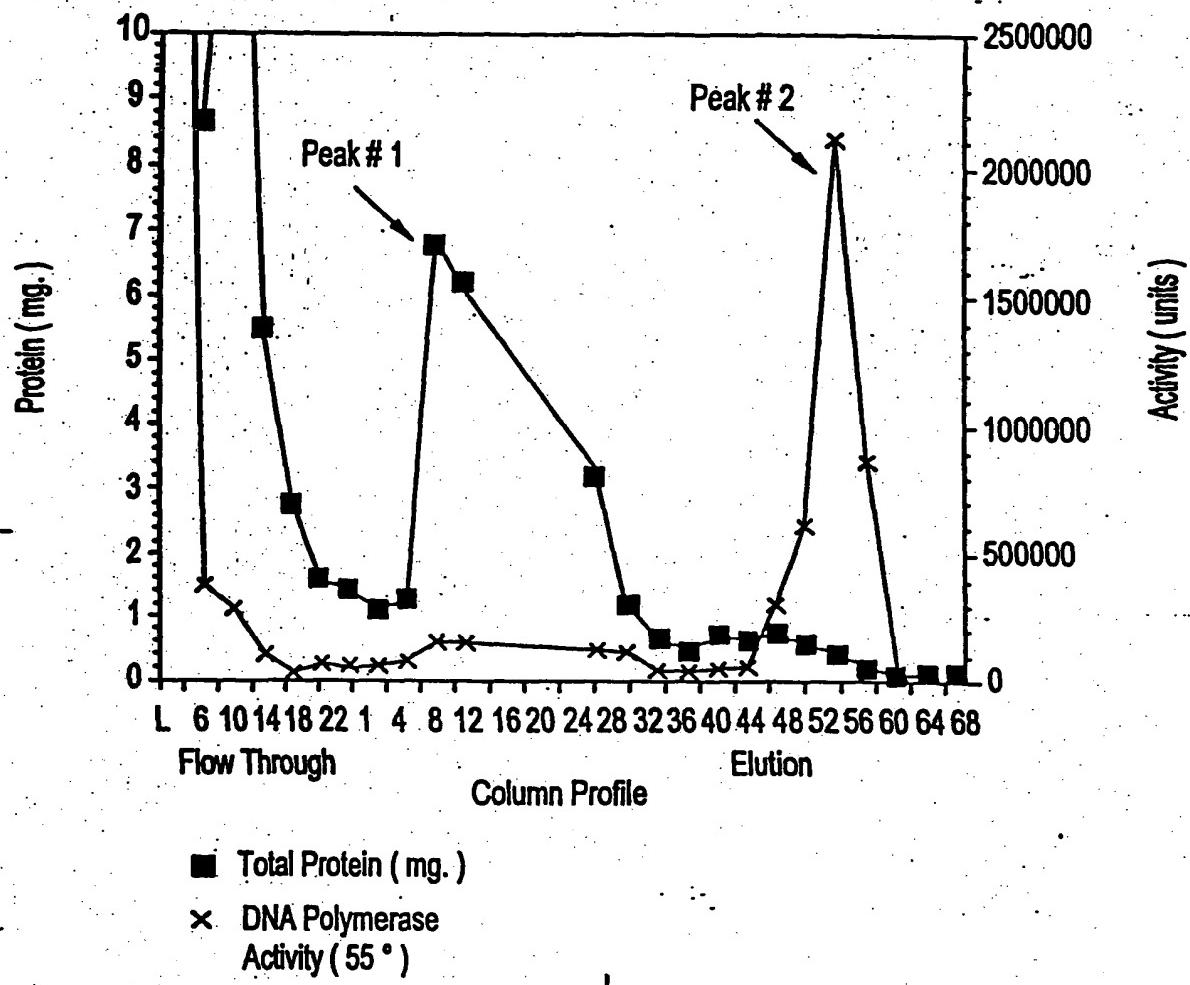


FIG.13B

ATP Agarose Step Column

FIG. 13C

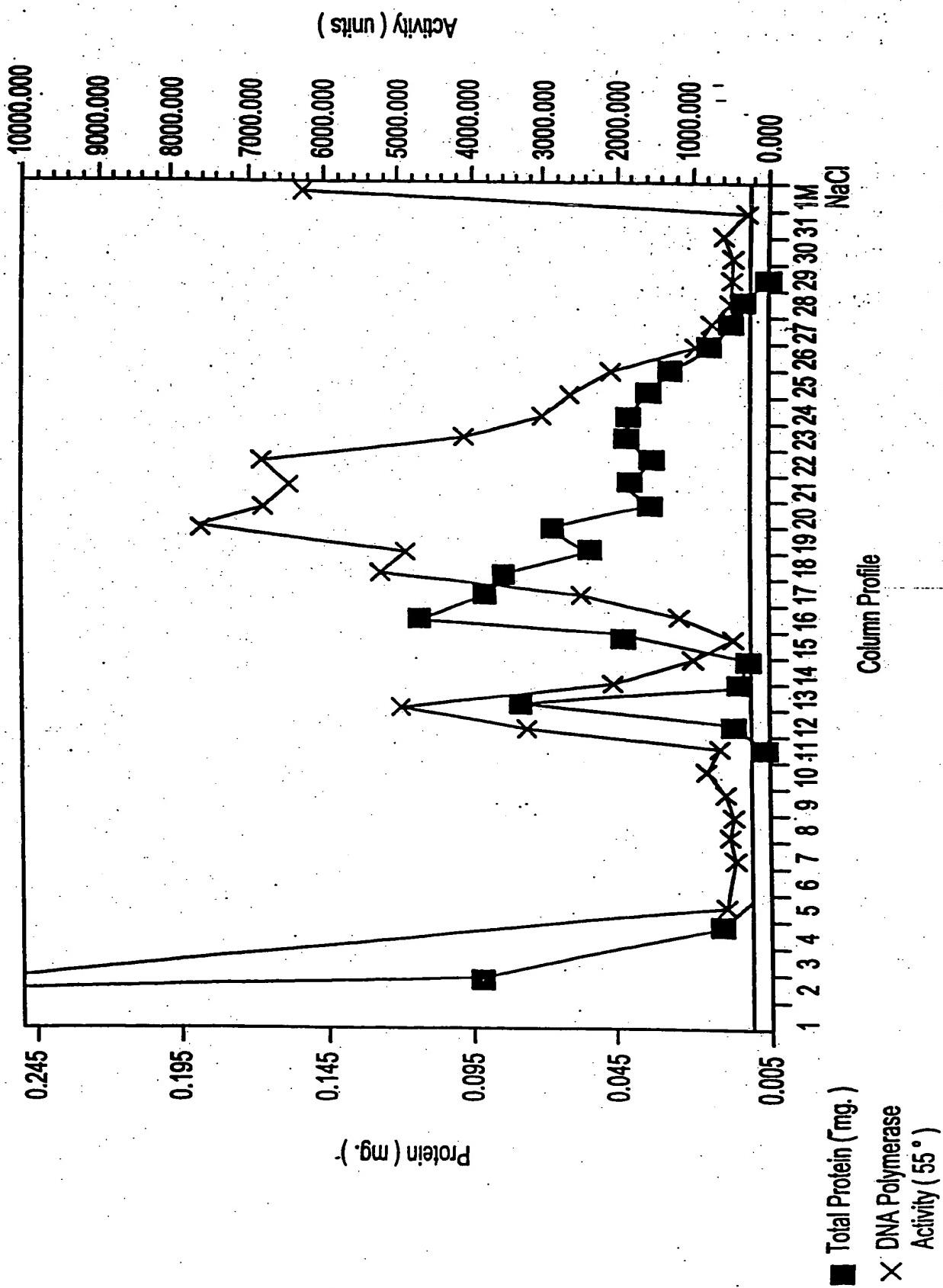


FIG. 14A

load FT 9 10 11 12 13 14 15 16 17 18 19 E. coli
 $\alpha\gamma\delta$

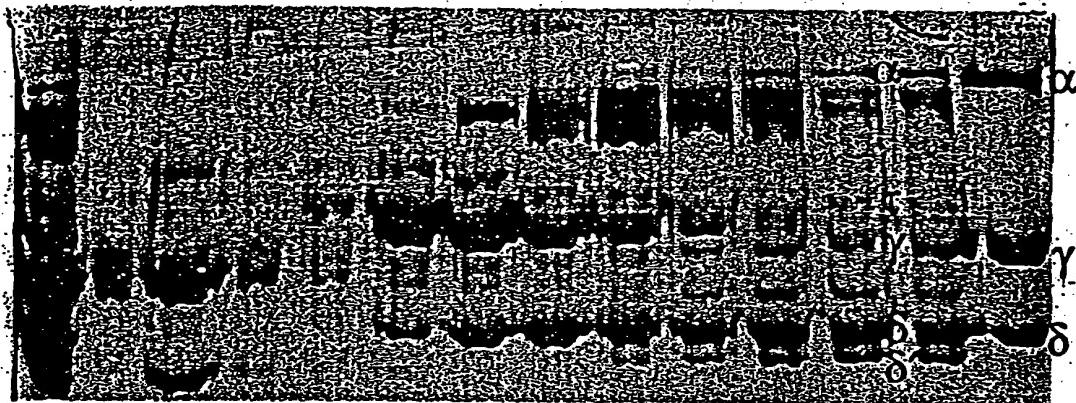


FIG. 14B

load FT 9 10 11 12 13 14 15 16 17 18 19



Alignment of Tth1 with alphas subunits of other organisms.

E. coli	DRYFLELIRTRGRADEESYIHLAAVELAEARGLPVV	197	(ID#72)
V. chol.	DHFYFLALSRRTGRPNEERYIQAALKLAERCDLPLV	197	(ID#73)
H. inf.	DHFYFLALSRRTGRPNEERYIQAALKLAERCDLPLV	197	(ID#74)
R. prow.	DRFYFEIMRHDILPEEQFIENSYTQIASELSIPIV	195	(ID#75)
H. pyl.	DDFYLEIMRHGILDDQRFTIDEQVTKMSLETGLKII	213	(ID#76)
S. sp.	DDYYLEIQDHGSVEDRLVNINLVKIAQELDIKIV	202	(ID#77)
M. tub.	DNYFLEIIMDHGLTIERVRDGLLEIGRALNIPPL	220	(ID#78)
T. th.	FFIEIIONHGLSEQK		(ID#61)

FIG. 15A

Alignment of Tth2 with alphas subunits of other organisms.

E. coli	NKRRAKNGEPPLDIAAIPLDDKKSFDMLQQRSETTAVFQLESRGMKD	618	(ID#79)
V. chol.	NPRLKKAGKPVRIEAIPLDDARSFRNLQDAKTTAVFQLESRGMKKE	618	(ID#80)
H. inf.	NVRMVRREGKPRVDIAAIPLDDPESFELLKRSETTAVFQLESRGMKD	618	(ID#81)
R. prow.	CKKLKEQGIKIDFDDMTEDDKKTYQMLICKGKGVGVFQFESTIGMKD	624	(ID#82)
H. pyl.	LKIITQHKISVDFLSLDMDDPKVYKTIQSGDTVGIFQIES-GMFQ	648	(ID#83)
S. sp.	QERKALQIRARTGSKKLPPDDVKKTHKLLLEGIFQLESQGMKQ	643	(ID#84)
M. tub.	IDNVVRAMRGIDLDDLESVPILDVKATYELLGRGDTTLGVFQLDGGPMRD	646	(ID#85)
T. th.	RVELDYDALTLDD		(ID#60)

FIG. 15B

ATGGGCCGGGAGCTCCGCTCGCCCACCTCCACCAGCACA	
CCCAGTTCTCCCTCTGGACGGGGCGCGAAGCTTCCGA	120
CCTCCTCAAGTGGGTCAAGGAGACGACCCCCGAGGACCCC	
GCCTTGGCCATGACCGACACGGCAACCTCTCGGGGCCG	
TGGAGTTCTACAAGAACGCCACCGAAATGGGCATCAAGCC	
CATCCTGGCTACGAGGCCTACGTGGCGGGAAAGCCG	240
TTTGACCGCAAGCGGGAAAGGGCTAGACGGGGCTACT	
TTCACCTCACCCCTCTCGCCAAGGACTTCACGGGGTACCA	
GAACCTGGTGCGCCCTGGCGAGCCGGCTTACCTGGAGGGG	360
TTTACGAAAAGCCCCGGATTGACCGGGAGATCCTGCGCG	
AGCACGCCGAGGGCCTCATGCCCTCTCGGGGTGCCCTCGG	
GGCGGAGATCCCCCAGTTCATCCTCCAGGACCGTCTGGAC	480
CTGGCCGAGGCCGGCTAACGAGTACCTCTCCATCTTCA	
AGGACCGCTTCTCATCGAGATCCAGAACCAACGGCCTCCC	
CGAGCAGAAAAAGGTCAACGAGGTCTCAAGGAGTTGCC	600
CGAAAGTACGGCTGGGATGGTGCCACCAACGACGGCC	
ATTACGTGAGGAAGGAGGACGCCCGCCACGAGGTCT	
CCTCGCATCCAGTCCAAGAGCACCCCTGGACGACCCGGG	720
CGCTGGCGCTTCCCTGCGACGAGTTCTACGTGAAGACCC	
CCGAGGAGATGCCGGCATGTTCCCCGAGGAGGAGTGGGG	
GGACGAGCCCTTGACAACACCCTGGAGATCGCCGCATG	840
TGCAACGTGGAGCTGCCCATGGGACAAGATGGTCTACC	
GAATCCCCGCTTCCCCCTCCCCGAGGGCGGACCGAGGC	
CCAGTACCTCATGGAGCTCACCTCAAGGGCTCCTCCGC	960
CGCTACCCGGACCGGATCACCGAGGGCTTCTACCGGGAGG	
TCTTCCGCCTTTGGGAAGCTTCCCCCCCACGGGACGG	
GGAGGCCCTGGCCGAGGCCCTGGCCCAGGTGGAGCGGGAG	1080
GCTTGGGAGAGGCTATGAAGAGCCTCCCCCTTGGCCG	
GGGTCAAGGAGTGGACGGCGGAGGCCATTTCCACCGGGC	
CCTTACGAGCTTCCGTATAGAGCGATGGGTTTCCC	1200
GGCTACTCCTCATCGTCAGGACTACATCAACTGGGCC	
GGAGAAACGGCGTCTCCGTGGGCCGGCAGGGGAGCGC	
CGCCGGGAGCCTGGTGGCCTACGCCGTGGGATACCAAC	1320
ATTGACCCCTCCGCTTCCGCTCTTGGCGCTTCC	
TGAACCCGGAGAGGGCTCCATGCCGACATTGACACGGA	
CTTCTCCGACCGGGAGCGGGACCGGGTATCCAGTACGTG	1440
CGGGAGCGCTACGGCGAGGACAAGGTGGCCAGATCGGA	
CCCTGGGAAGCCTCGCCTCAAGGCCGCCCTCAAGGACGT	
GGCCCGGGTCTACGGCATCCCCACAAGAACGGGAGGAA	1560
TTGGCCAAGCTCATCCGGTGCAGTTGGGAAGCCCAAGC	
CCCTGCAGGAGGCCATCCAGGTGGTGGAGCTTAGGGC	
GGAGATGGAGAACCGCCAAAGGTGCGGGAGGTCTCGAG	1680
GTGGCCATGCGCTGGAGGGCCTGAACCGCCACGCCCTCG	
TCCACGCCGCCGGGTGGTATCGCCGCCAGCCCCCTCAC	
GGACCTCGTCCCCCTCATGCGCACCAGGAAGGGCGGCC	
GTCACCCAGTACGACATGGGGCGGTGGAGGCCTTGGGGC	1800
TTTGAAAGATGGACTTTTGGGCCCTCGCACCCACCTT	

FIG. 16A

CCTGGACGAGGTCAAGCGCATCGTCAAGGCCTCCAGGGG	1920
GTGGAGCTGGACTACGATGCCCTCCCCCTGGACGACCCA	
AGACCTTCGCCCTCCTCTCCGGGGGAGACCAAGGGGGT	
CTTCCAGCTGGAGTCGGGGGGATGACCGCCACGCTCCGC	
GGCCTCAAGCCGGCGCTTGAGGACCTGATGCCATCC	
TCTCCCTCTACCGCCCCGGGCCATGGAGCACATCCCCAC	
CTACATCCGCCGCCACCACGGCTGGAGCCCCTGAGCTAC	
AGCGAGTTTCCCCACGCCAGAAGTACCTAAAGCCCATCC	
TGGACGAGACCTACGGCATCCCCGTCTACCAGGAGCAGAT	
CATGCAGATCGCCTCGGCCGTGGCGGGTACTCCCTGGGC	
GAGGCGGACCTCCTGCCGGGTCCATGGGCAAGAAGAAGG	
TGGAGGAGATGAAGTCCCACCGGGAGCGCTCGTCCAGGG	
GGCCAAGGAAAGGGCGTCCCCGAGGAGGAGGCCACCGC	
CTCTTGACATGCTGGAGCCTCGCCAACCTACGGCTTCA	
ACAAATCCCACGCTGCCGCTACAGCCTCCTCTCCCTACCA	
GACCGCCTACGTGAAGGCCACTACCCGTGGAGTCATG	
GCCGCCCTCCTCTCCGTGGAGCGGCCACGACTCCGACAAGG	
TGGCCGAGTACATCCCGCACGCCGGGCACTGGGATAGA	
GGTCCTTCCCCGGACGTCAACCGCTCCGGGTTGACTTC	
CTGGTCCAGGGCCGGCAGATCCTTTCGGCCTCTCCGCGG	
TGAAGAACGTGGCGAGGCCGGGGGGGAGGCCATTCTCCG	
GGAGCGGGAGCGGGGGGGCCCTACCGGAGCCTCGCGAC	
TTCCTCAAGCGGCTGGACCGAGAACGGTGCCTAACAGCGGA	
CCCTGGAGTCCCTCATCAAGGCGGCCCTGGACGGCTT	
CGGGGAAAGGGCGCGCTCCTCGCCTCCCTGGAAAGGGCTC	
CTCAAGTGGCGGCCGAGAACCGGGAGAACGCCGCTCGG	
GCATGATGGGCCTCTCAGCGAACGGAGGCCGCTT	
GGCCGAGGCCGCCCTGGACGAGATCACCGGCTCCGC	
TACGAGAACGGAGGCCCTGGGATCTACGTCTCCGGCCACC	
CCATCTTGGGTACCCGGCTCCGGGAGACGGCCACCTG	
CACCCGGAGGGAGCTCCCGACCTGGCCCGGGACCTGCCG	
CCCCGGTCTAGGGTCTCCTGGGGAGGGAGTGGTGGAGGAGG	
TGGTGCAGAACGCCACAAAGAGCGGGGGAGTGTGGCCCG	
CTTCGTCTCTCCGACGAGACGGGGCGCTTGAGGCGGTG	
GCATTGGCCGGGCTACGACCGAGGTCTCCCGAGGCTCA	
AGGAGGACACCCCGTGCTCGCCTCGCGAGGTGGAGCG	
GGAGGAGGGGGCGTGGGGTGTGGCCCAGGCCGTTGG	
ACCTACGAGGAGCTGGAGCAGGTCCCCGGGCCCTCGAGG	
TGGAGGTTGGAGGCCTCCCTGGACGACCGGGGGGTGGC	
CCACCTGAAAAGCTCTGGACGAGCACGCCGGGACCCCTC	
CCCCTGTACGTCCGGTCCAGGGCGCCTCGCGAGGCC	
TCCTCGCCCTGAGGGAGGTGGGGGGAGGAGGAGGCTGT	
AGGCAGGCCGCGTGGTCCGGGCTACCTCTGCCGACCG	
GGAGGTCTTCTCCAGGGCGCAGGCAGGGGGAGGCCAG	
GAGGCGGTGCCCTCTAGGGGTGGGCCGTGAGACCTAGC	
GCCATCGTTCTGCCGGGGCAAGGAGGCCCTGGGCCGAC	
CCCTTTGG	

FIG. 16B

MGRELRF AHLHQHTQFSLLDGAPKLS DLLKWVEETTPEDP	
ALAMTDHGNLFGAVEFYKKATEMGIKPILGYEAYVAAESR	120
FDRKRGKGLGGYFHLTLAKDFTGYQNLVRLASRAYLEG	
FYEKP RIDREI LREHAEGLIALSGCLGAEIPQFILQDRLD	
LAEARLINEYLSIFKDRFFIEIQNHLPEQKKVNEVLKEFA	
RKYGLGMVATNDGHYVRKEDARAHEVLLAIQSKSTLDDPG	240
ALALPCEE FYVKTPEEMRAMFPEEEVGGRSPLTTPWRSRH	
VQRGAAIGTRWSTRIPRFPLPEGRTEAQYLMEFTFKGLLR	
RYPDRITEGFYREVFRSLSGKLPPHGDEALAEALAOVERE	360
AWERLMKSLPPLAGVKEWTAEEAIFHRALYELSAIERMGFP	
GLLPHRPGLHQLGPEKGVS VGPGRGGAAGSLVAYAVGITN	
IDPLRFGLLFERFLNPERVSMPDI DTFSDRERDRV IQYV	480
RERYGEDKVAQIGT LGS LASKA ALKEVARVYGI PRKKAEE	
LAKLI PVQFGKPKPLQEAIQVVPELRAEMEKDPKV REVLE	
VAMRLEG LNRHASVHAGRGGVFSEPLTDLVPLCATRKGGP	600
YTQYDMG AVEAL GLLKMDFLGLRTLTFLDEVKRIVKASQG	
VELDYDALPLDDPKTFALLSRGETKGVFQLES GGMTATLR	
GLKPRRFEDLIAILS LYRPGPMEH IPTYI RHHGLEPVSY	720
SEFPHAEKYLKPILDETYGIPVYQE QIMQIASAVAGYSLG	
EADLLR RSMGKKV EEMKSHRERFVQGAKERGVPEEEANR	
LFDMLEAFANYGF NKSHAAAYSLLSYQTAYVKAHYPVEFM	840
AALLSVERHDSDKVAEYIRDARAMGIEVLPDVNRSGFDF	
LVQGRQILFGLSAVN VGEAAA EAILRERERGGPYRSLGD	
FLKRLDEKVLNKRTLESLIKAGALDG FGERARLLASLEGL	960
LKWAAENREKARSGMMGLFSEVEEPLAEEAALDEITRLR	
YEKEALGIYVSGHPILRYPGLRETATCTLEELPHLARDLP	
PRSRVLLAGMVEEVVRKPTKSGGMMARFVLSDETGALEAV	1080
AFGRAYDQVSPRLKEDTPVLVLAEVEREEGGVRVLAQAVW	
TYQELEQVPRALEVEEA SLPDDR GVAHLKSLLDEHAGTL	
PLYVRVQGAFGE ALLALREV RVGEA LGALEAAGFPAYLL	
PNREVSPRLTGSGGPRGRALSTGLALKTYPIALPGGNEAL	
ARPLL	1200

FIG. 16C

	Start1	Start2	3' - EXO I	3' - EXO II	3' - EXO III
T. th.	VERVVRTILLDGRFLLEEGVGLWEWRYPFPLEGEAVVVLDILETTGLAG	-----LDEVIEGLLRLEGG	-----RRLPF	-----QSLVR-PLPP--AEARSWNLT--GIPREALEEAPSLEEVELEKAYPLRGDDATLVTHNAAFDLGFL-RPALEGGLG	-----YRLENPVVDLRLARRGLPGLRRYGLDAISSEVLELPRT--CHRALEEDVERTLAVVHEVYMLT-----SG
D. rad.	PWPQDVVVFDILETTGFSPA	-----SAAIVEIGAVRIVGGQIDETILKF		ETLVR-PTRPDGSMILSIPWQAQRVHGISDEMVRAPAXKDVL.PDFDFDVDSAVVAENVSEDGGM-RAGAERLG	-----LSWAPERELCTMQLSRRAFPRTHTNLTVIAERLGLEFAPGGRHSYGDVQVTAQAYLRLIELIG-----ER
Bac. sub.	HGIKMIYGMEANLVDDGVPPIAYNAAHRILLEEEETXVVFDVETTGLSAV	-----YDTIIELAAVKVKGGE--TIDKF		EAFAN-PHRP--LSATIIELT--GITDDMLQDAPDVWDVIRDFREWIGDDILVAAHNASFTDMGFL-NVAYKKLL	E---VEKAKNPVTDLQMARQMPGKRN-NLDALCDRLGIDNSKRTLLEGALLDAEILADVYLMMTGGOTNLFDEEE
H. inf.	MINPNRQRIVLDFETTGMMQLGAHYEGHCIEIGAVELINRR-YTGNNX			H.IYIK-PDRP--XDPDAIKVH--GITDEMILADKPEFKEVAQDFLDYINGAELLIENAPFDVGFM-DYEFRKLK	-LNVKTDDICLVTDTLQMARQMPGKRN-SLDALCARYEIDNSKRTLLEGALLDAQILAEVYLAMTGGOTSMAFAME
E. c.	MSITAITRQTVLDFETTGMMQIGAHSEGKIIIEIGADEVVNRRL-TGNNF			HVYLK-DRLV---DPEAFGVH---GIAVDFLLDKPTFAEVAVEFMDYTRGAELEVTHNAAFDIGM-DYEFSLLK	RDIAKTNTFCKVTDLSLAVARKMFPGKRN-SLSFLKAILSRY-SLSFLK
H. pyl.	NLEYLKACGLNFETSENLTKLKNLKTPLKDEVFSPIIDLETTGSCPI	-----KHEITLEIGAVQVRGGE--TINRF		ETLVVKVSVP---DYTAELT---GITYEDTLNAPSAAHEALQEIRLFLGNSVFTAHHANFDYNFLGRYFVEKLH	-----CPLLNLKLCFLDSKRAILSRY-SLSFLK

FIG. 17

FIG. 18A

ATGGTGGAGCGGGTGGTGCGGACCCCTCTGGACGGGAGGT 40
TCCTCCTGGAGGAGGGGGTGGGGCTTGGGAGTGGCGCTA
CCCCTTCCCTGGAGGGGGAGGCCTGGTGGTCCCTGGAC 120
CTGGAGACCACGGGGCTTGCCGGCCTGGACGAGGTGATTG
AGGTGGGCCTCCTCCGCCTGGAGGGGGGAGGCCTCCC 200
CTTCCAGAGCCTCGTCCGCCCTCCGCCCGCCGAAGCC
CGTCGTGAAACCTCACCGGATCCCCCGGAGGCCCTGG 280
AGGAGGCCCTCCCTGGAGGAGGTTCTGGAGAAGGCCTA
CCCCCTCCGCCGACGCCACCTGGTATCCACAACGCC 360
GCCTTGACCTGGCTTCCTCCGCCGCCCTGGAGGGCC
TGGGCTACCGCCTGGAAAACCCGTGGACTCCCTGCG 440
CTTGGCCAGACGGGGCTTACCAAGGCCTTAGGCCTACGGC
CTGGACGCCCTCTCCGAGGTCTGGAGCTTCCCGAAGGA 520
CCTGCCACCGGGCCCTCGAGGACGTGGAGCGCACCCCTCGC
CGTGGTGCACGAGGTATACTATATGCTTACGTCCGGCCGT 600
CCCCGACGCTTGGAACTCGGGAGGTAG

MVERVVRTLLDGRFLLEEGVGLWEWRYPFPLEGEAVVLD 40
LETTGLAGLDEVIEVGLLRLEGGRRLPFQSLVRPLPPAEA
RSWNLTGIPREALEEAPSLEEVLKAYPLRGDATLVIHNA 120
AFDLGFLRPALEGGLGYRLENPVVDSLRLARRGLPGLRRYG
LDALSEVLELPRTCHRALEDVERTLAVVHEVYYMLTSGR 200
PRTLWELGRZ

FIG. 18B

Alignment of dnaA genes.

P. mar.	MLEASWEK VQSSL--KQNLSK--	-----PSYE TWIRPTEESG--FKN CELTTLAPNSFSSAW LKQNYSQTIQETAE-		
Syn. sp.	MVSCENLWQQ ALAII--ATQLTK--	-----PAFD TWIKAVSLIS--LCD GVTIQQVENGVLNH LQKSYGLIMEVLT-	65	65
B. sut.	MENILDLWNQ ALAQI--ECKLSK--	-----PSFE TMIKSTKAHS--LQC DLTITAPNEFARDW LESRYLHLIAIDTY-	67	67
M. tub.	MTDDPGSGFTTWNA WSELNGDPKVDDGP	SSDANI SAPLTPQQR AWIUNLVQELT--IVE GFALLSVPSSEFVQNE TERHRAPITDALS-	67	67
T. th.	MSHEAWOIH VLEHI--RRSITE--	-----VEFH TIFERIRPLG--IRD GVELEAVPTSEALDW IRRHYAGLIQEGRP-	87	87
E. coli	MSTSLIWQQ CLARL--QDELPA--	-----TEFS MMRPLQAE--LSD NTLLAYAPNREFLDW VRDKYLYANNINGLT-	66	66
T. mar.	MRER ILQEI--KTRVNR--	-----KSWE LMFSSFDWRS--IBG NKVVFSGNLFIIKEW LEKKYYSVLSKAVK-	61	61
H. pyl.	MDTNNNIEKE ILALKQNPKVSL--	-----IEYE NYTSOLKVNPNASKS DIAFFYAPNQVLT ITARYGALLEKIELSQ	72	72
P. mar.	EIFG----EPVTVHK VKNAEESSEDEHYSSA P-----	-----ITPPILEASPGSV DSSGSSSLRSLK-----	130	130
Syn. sp.	DLTG----QETIVKLI TDGLEPHS---LICQ E-----	-----SSLPMETTP-----	115	115
B. sut.	ELTG----EELS IKFV IPONODVEDEMPKPQ	VKKAVKEDTSDEPON -----	119	119
M. tub.	RLIGH-QIQLGVRLA PPATDEADDTTVPPS	ENPATSPDTITND EIDDSAAARGDNQHS WPSYTFTERPHNTDSA TAGTTSLNRRYTFDT	119	119
T. th.	LIGAQ-AFRFELRVW PGWWQEDIFOPPPS	PPQAQQP-----	176	176
E. coli	SFCGADAPOLRFEG TKPVTQTPOAATSN	VAAPAOVQAQTOPORA APSTRSGMDNVPA PA EP-----	108	108
T. mar.	VVLG----NDATFETT YEAFEPHSSTYSEPLV	KKRAVLLTP-----	140	140
H. pyl.	NKVVG-MHIAHSVDVR IEVAKPQIQNAQSNI	NYKAIKTS-----	106	106
		-----VKSQSYTFEN	118	118
P. mar.	FVVGPNSRMAAHAAAM AVAESPGREFNPFLI CCGVGLGKTHLMQAI GHYRLEIDPGAKVSY VSTETTFNDLIL--A	IRDQRMQAERDRYR-	217	217
Syn. sp.	FVVGPNTNMRMAHAASL AVAESPGREFNPFLI CGGVGLGKTHLMQAI AHTRLEMVNAKVVY VSTERFTNDLIT--A	IRDQNMEDFRSYR-	202	202
B. sut.	FVIGSGNRFHAASL AVAEAPAKAYNPFLI YGGVGLGKTHLMHAI GHYVTDHMPSAKVVY LSSEKFTNEFIN--S	IRDANKAVDFNRYR-	206	206
M. tub.	FVIGASNRFHAAAAL AIAEAAPARAYNPFLI WCESGLGKTHLHAA GENYAQRLLPGMRVKY VSTEEFTNDFIN--S	LRDRKVAFKRSYR-	263	263
T. th.	SMWGPTTPWPHGGAV AVAESPGRAVNPFLI YGGRGLGKTYMHAV GPLRAKREPMEARLEY VSTEEFTNELINRPS AR-DRMTEFRYR-	196	196	
E. coli	FVEGKSNQLARAAR QVADNPGGAYNPFLI YGGTGLGKTHLHAV GNGTMARKPAKVVY MHSERFTVDMVK--A LQNNATEEFKRYR-	227	227	
T. mar.	FVVGPGENSFAYHAL EVAKHPGR-YNPFLI YGGVGLGKTHLQSI GNYWVQNEPDLRVMY ITSEKFLNDLV--S MKEGKLNEFREKRYK	193	193	
H. pyl.	FVVGSCTNNTYEIAK KVAQSDTIPPYNPVLF YGGTGLGKTHLMNAI GNAHALEK--HKKVL VTSEDFTLDFLK--H LQNKTMDSFKAKYR-	203	203	

FIG. 19A

FIG. 19B

P.mar.	AADLILVDDIOFIEG KEYTQEEFFFTFNL HDAGSQIVLASSDRPP	SQIPRLQERUMSRFS MGLIADVQAPDLETR MAJLQKKAEEHVRGL	307
Syn. sp.	SADFLILLDDIOFIEG KEYTQEEFFFTFNL HEAGKQVWVASDRP	QRIPLQDRLLISRF S MGLIADVQAPDLETR MAJLQKKAEEHVRGL	292
B.sut.	NVDVLILLDDIOFIEG KEQIQEEFFFTFNL HEESKQIVTSSDRPP	KELPTLEDRLRSRF E MGLITDITPPDLETR IAIILRKKAQMERLDI	296
M.tub.	DVDVLILLDDIOFIEG KEGIQEEFFFTFNL HNANKQIVTSSDRPP	KOLATLEDRLRTRE E MGLITDQQPDETR IAIILRKKAQMERLAV	353
T.th.	SVDLILLDDIOFIEG KERTQEEFFFTFNL YEAKHQITSSDRPP	KOLITTLEAULRSRF E MGLITDNPADLETR IAIILRKKAQMERLAV	285
E.coli	SVDALILLDDIOFIEG KERSQEEFFFTFNL LEGNQITLTSDRPP	KEINGVEDRLKSREG E MGLITVAEPELETR VALIMKKADENDIRL	317
T.mar.	KVDILLILLDVQFLIG KTGVQTELFITFNL HDSGCKQIVICSDREP	QKLSEFODLVSRQ MGLVAKLEPPDEETR KSTARKTLEIERGEL	283
H.py1.	HCDFFLIDDAQFLQG KPKLKEEFITFNL HANSKQIVLISDRSP	KMLAGEDRLKSRF E MGLITAKUMPPDLETR LSVVKQRCQLNQITL	293
P.mar.	PRDLIQFTAGRETSN IRELEGALITRAIAFYA SITGCPMTVDSTAPM	LD----PNCQGVENT PKQVLDKVAEVTKT PDEMRSASRRR-FVS	392
Syn. sp.	PKEVIEYLASHYTSN IRELEGALIRAYT SLNVANTVENLAFV	LN----PPVEKVAAP PETLITTVAQHYQLK VEELLLNSRRR-EVS	377
B.sut.	PNEVMLYIANQIDSN IRELEGALIRWAYS SLINKDINADAAEA	LKDII-PSSSPKPVIT IKEIQRUVGQOFNIK LEDFKAKKRKT-SVA	384
M.tub.	PDVYLELIASSERN IRELEGALIRVTAFA SLANKTPIDKALAETV	LRDLI-ADANTMOIS AATIMAATAEYFTT VEELLRGPCKTR-ALA	441
T.th.	PEDALEYIARQVTSN IREMEGALMRASPFA SLNGVELTRAVAAKA	LRHLR-P--RELEAD PLEIRKAAGPVRPE TPGGAHGERRKKEVV	372
E.coli	PGEVAFFIAKRLRSN VRELEGALMRVIANA NFGRAITTDFUREA	LRDLI-A-LQEKLVT IDNIQRTVAEYKIK VADLLSKRSR-SVA	404
T.mar.	PEEVLNFTVAENVDDN IRRRGALIKLIVK ETIGKEVDLKAEATLL	IKOFIKPNRKAMDP IDELIEVAKVTPGP REETLNSNSRNV-KAL	372
H.py1.	PEEVMEYLAQHISDN IRQMEGALIKISVNA NMNMASTDILNLAKTV	LEDL--QKDHAEGSS LENTLAVAQSINK SSEIKVSSSRQK-NVA	380
P.mar.	QARQVGMYLMRQGTN LSLPRIGDTFGGKDH TTVMYAJEQVEKLS	S-----DPQIA SQVQKIRDLQIDSR RKR----	461
Syn. sp.	LARQVGMYLMRQHTD LSLPRIGEAFGGKDH TTVMYSCDKITQOQQ	K-----DMETS QTLTSLSRINTIAQG APES---	447
B.sut.	FPRQIAMYLSREMID SSLPKIGEEFGGRDH TTVAHAHEKISKLLA	D-----DEQLQ QKVKIKEQQLK-----	446
M.tub.	QSRQIAMYLCRELTID LSLPKIGQAFG-RDH TTVMYAQRKILSEMA	E-----RREVF DVKEKLTTRIQRSK R-----	507
T.th.	LPRQIAMYLVRELTP ASLPEIGQLEFGGRDH TTVMYAJQKVQELAG	KP-----DREVO GLLRTLREACTDPVD NLWTTCG	446
E.coli	RPRQTMAMALAKELTN HSLPEIGDAFGGRDH TTVLHACRKIEQLRE	E-----SHDIK EDFSNMLRTLSS-----	467
T.mar.	TARRIGMYVAKNYK SSLRTIAEKFN-RSH FVVDVSKVKDSIL	KG-----NKQLK ALIDVEGEISRAL SG-----	440
H.py1.	LARKLWVFARLYTP NPTLSIAQFLDQKDH SSISRMYSGVKKMLE	EKSPFPVLSRREEIK NRNLNEENDKTAFNS SE-----	457

GTGTCGACGAGGCCGTGGAACACAGTTCTGGAGCACA
TCCGCCGCAGCATCACCGAGGTGGAGTCCACACACTGGTT
TGAAAGGATCCGCCCTTGGGATCCGGACGGGTGCTG 120
GAGCTGCCGTGCCACCTCCTTGGCCCTGGACTGGATCC
GGCGCCACTACGCCGGCTCATCCAGGAGGGCCCTCGGCT
CCTCGGGGCCAGGCAGCCCGGTTGAGCTCCGGTGGTG 240
CCCAGGGTCGTAGTCAGGAGGACATCTCCAGCCCCCGC
CGAGCCCCCGGCCAAGCTAACCGAAGATAACCTTAA
AACTTCGTGGTGGGCCAACAACTCCATGGCCCCACGGC 360
GGCGCCGTGGCCGTGGCGAGTCCCCCGGCCGGCCTACA
ACCCCCCTTCATCTACGGGGCCGTGGCCTGGAAAGAC
CTACCTGATGCACGCCGTGGCCACTCCGTGCGAACGCG 480
TTCCCCCACATGAGATTAGAGTACGTTCCACGGAAACTT
TCACCAACGAGCTCATCAACCGGCATCCGCGAGGGACCG
GATGACGGAGTTCCGGAGCGGTACCGCTCCGTGGACCTC 600
CTGCTGGTGGACGACGTCCAGTTCATGCCGGAAAGGAGC
GCACCCAGGAGGAGTTTCCACACCTCAACGCCCTTA
CGAGGCCACAAGCAGATCATCCTCTCCGACC GGCG 720
CCCAAGGACATCCTCACCCCTGGAGGCGCGCTGCGGAGCC
GCTTGAGTGGGGCTGATCACCGACAATCCAGCCCCCGA
CCTGGAAACCGGATGCCATCCTGAAGATGAACGCCAGC 840
AGCGGGCCTGAGGATCCCAGGACGCCCTGGAGTACATCG
CCCGGCAGGTACCTCCAACATCCGGAGTGGAAAGGGGC
CCTCATGCGGGCATGCCCTTCGCCCTCCCTAACGGCGTT 960
GAGCTGACCCCGCCGTGGCGGCCAAGGCTCTCCGACATC
TTCGCCCCAGGGAGCTGGAGGCGGACCCCTGGAGATCAT
CCGCAAAGCGCGGGACCAAGTTCGGCCTGAAACCCGGGA 1080
GGAGCTCACGGGAGCGCCGCAAGAAGGAGGTGGTCTCC
CCCGGCAGCTGCCATGTACCTGGTGCAGGGAGCTCACCC
GGCCTCCCTGCCGAGATCGACCAGCTAACGACGACCGG 1200
GACCACACCACGGTCTCTACGCCATCCAGAAGGTCCAGG
AGCTCGCGAAAGCGACCGGGAGGTGCAGGGCCTCCTCCG
CACCCCTCCGGGAGGCCTGCACATGA

FIG.20A

VSHEAVWQHVLEHIRRSITEVEFHTWFERIRPLGIRDGV
ELAVPTSFALDWIRRHYAGLIQEGRLLGAQAPRFELRVV
PGVVVQEDIFQPPPSPPAQAPEDTFKTSWWGPTTPWPHG 120
GAVAVAESPGRAYNPLFIYGGRGLGKTYLMHAVGPLRAKR
FPHMRLEYVSTETFTNELINRPSARDRMTEFRERYRSVDL
LLVDDVQFIAGKERTQEFFHTFNALYEAHKQIILSSDRP 240
PKDILTLEARLRSRFEWGLITDNPAPDLETRIAILKMNAS
SGPEDPEDALEYIARQVTSNIREWEGLMRASPFAASLNGV
ELTRAVAAKALRHLRPRELEADPLEIIRKAAGPVRPETPG 360
GAHGERRKKEVVLPRQLAMYLVRELTPASLPEIDQLNDDR
DHTTVLYAIQKVQELAESDREVQGLLRTLREACT

FIG.20B

ATGAAACATAACGGTTCCCAAAA	40
ACTCCTCTCGGACCAGC	
TTTCCCTCCTGGAGCGCATCGTCCCCTCTAGAAGCGCAA	
CCCCCTCTACACCTACCTGGGGCTTACGCCGAGGAAGGG	120
GCCTTGATCCTCTCGGGACCAACGGGGAGGTGGACCTCG	
AGGTCCGCCCTCCCCGCCGAGGCCAAAGCCTTCCCCGGGT	200
GCTCGTCCCCGCCAGCCCTCTCCAGCTGGTGCAGGC	
CTTCCTGGGACCTCGTGGCCCTCGGCCTCGCCTCGGAGC	280
CGGGCCAGGGGGGGCAGCTGGAGCTCTCCTCCGGCGTT	
CCGCACCCGGCTCAGCCTGGCCCTGCCGAGGGCTACCCC	360
GAGCTTCTGGTGCCCGAGGGGGAGGACAAGGGGCCTTCC	
CCCTCCGGACGCGGATGCCCTCCGGGAGCTCGTCAAGGC	440
CTTGACCCACGTGGCTACGCCGAGCAACGAGGGAGTAC	
CGGGCCATCTTCCGGGGGTGCAGCTGGAGTTCTCCCCCC	520
AGGGCTTCCGGGCGGTGGCCTCCGACGGTACCGCCTCGG	
CCTCTACGACCTGCCCTGCCCAAGGGTTCCAGGCCAAG	600
GCCGTGGTCCCCGCCGGAGCGTGGACGAGATGGTGCAGG	
TCCTGAAGGGGGCGGACGGGGCCGAGGCCGTCCCGCCCT	680
GGCGAGGGGGTGTGGCCCTGGCCCTCGAGGGCGGAAGC	
GGGGTCCGGATGGCCCTCCGCCTCATGGAAGGGGAGTTCC	760
CCGACTACCAGAGGGTCATCCCCCAGGAGTTGCCCTCAA	
GGTCCAGGTGGAGGGGAGGCCCTCAGGGAGGCCGTGCGC	840
CGGGTGAGCGTCCTCTCCGACCGCAGAACCAACCGGGTGG	
ACCTCCTTTGGAGGAAGGCCGGATCCTCCTCTCCGCCGA	920
GGGGGACTACGGCAAGGGCAGGAGGAGGTGCCCGCCAG	
GTGGAGGGGCCGGACATGGCCGTGGCCTACAAACGCCGCT	1000
ACCTCCTCGAGGCCCTCGCCCCCGTGGGGGACCGGGCCA	
CCTGGGCATCTCCGGGCCACGAGCCCAGCCTCATCTGG	1080
GGGGACGGGGAGGGTACCGGGCGGTGGTGGTGCCTCA	
GGGTCTAG	1128

FIG.21A

MNITVPKKLLSDQLSLERIVPSRSANPLYTYLGLYAE^G 40
ALILFGTNGEVDLEVRLPAEAQSLPRVLVPAQPFFQLVRS
LPGDLVALGLASEPGQGGQLELSSGRFRTRLSLAPAEGYP 120
- ELLVPEGEDKGAFPLRTRMPSGELVKALTHVRYAASNEEY
RAIFRGVQLEFSPQGFRAVASDGYRLALYDLPLPQGFQAK 200
AVVPARSVDEMVRVLKGADGAEAVLALGEGVLALALEGGS.
GVRMALRLMEGEFPDYQRVIHQEFALKVQVEGEALREAVR 280
RVSVLSDRQNHRVDLLLEEGRILLSAEGDYGKGQEEVPAQ
VEGPDMAVAYNARYLLEALAPVGDR AHLGISGPTSPSLI^W 360
GDGEGYRAVVVPLRVZ

FIG.21B

FIG. 22A

FIG. 22B

T. th. beta	GGSGVRLRUMEGEFPDYQRVITQEFALKVQVEGEALREAVRRVSVLSDRQNMHRVDLL	
E.coli.bet	---DFIFTTSKLVLDGRFPDYRRLPKNPDKHLEAGCDILKQAFARAAILSNEKFRGTRLYV	
P.mirab.be	---DFIFTTSKLVLDGRFPDYRRLPKNPDKHLEAGCDILKQAFARAAILSNEKFRGTRINL	
H.infl.bet	---NTVFTTSKLVLDGRFPDYRRLPKRNATKIVEGNWMLKQAFARASILSNERARSVRLSL	
P.put.beta	---EFTFTTSKLVLDGRFPDYRVLPKGDKLVVGDROALREAFSRTAILSNEKYRGIRQLQ	
B.cap.beta	---NLIFITQLIEGEYPDYSVLFKEKKNPITNSIIILKKSSLRVALLAHEKFCGIEIKI	
* * * * *		
T. th. beta	EEGRILLSAEGDYGK-GQEEVPAQVEGPDMAVAYNARYILLEALAPVG-DRAHLGIGSGPTS	
E.coli.bet	SENQLKITANNNPEQEEAEELIDVTYSGAEMEIGFNVSYLVDLNALKCENVRMCLTDVS	
P.mirab.be	TNGQLKITANNNPEQEEAEELIDVTYQYQGEEMEIGFNVSYLVDLNALKCENVRMCLTDVS	
H.infl.bet	KENQLKITASNNTEHEEAETTVDVNVNGEELEVEGENVTYILDVNLALKCNQVRMCLTDVS	
P.put.beta	AAGQLKIQANNNPEQEEAEEEEISVYEGSSLEIGFNVSYLVDLGWMTTEQVRLLISDSNS	
B.cap.beta	ENGKFKVLSNDNQEEETAEDLFEDIYFGEKIEISINVYLLDVINNIKSENIALFLNSKS	
* * * * *		
T. th. beta	PSLIWGDG-EGYRAVWPLRVZ	(ID#108)
E.coli.bet	SVQIEDAAQSAAAYVMPMRLZ	(ID#109)
P.mirab.be	SVQVENVASAAAAYVMPMRL-	(ID#110)
H.infl.bet	SCLLIENCEDSSCEYVTPMPRL-	(ID#111)
P.put.beta	SALIQEAGNDSSSYVMPMRL-	(ID#112)
B.cap.beta	SIQIEAENNSSNAYVMPMRL-	(ID#113)
* * * * *		

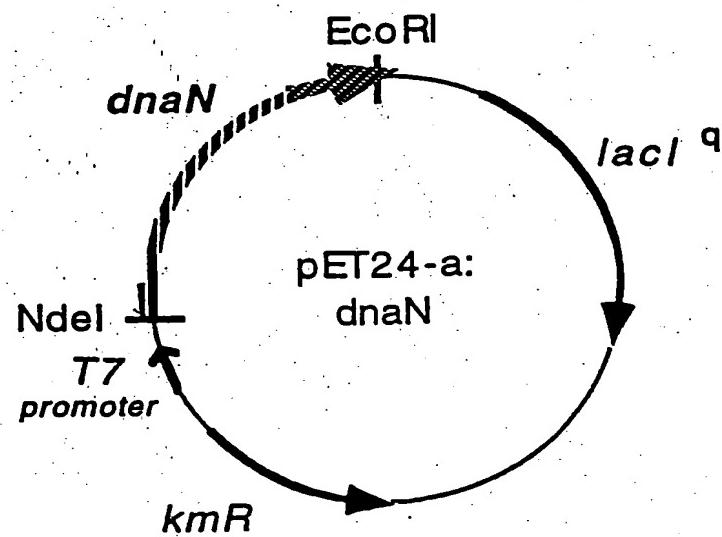


FIG.23

FIG. 24A Induction

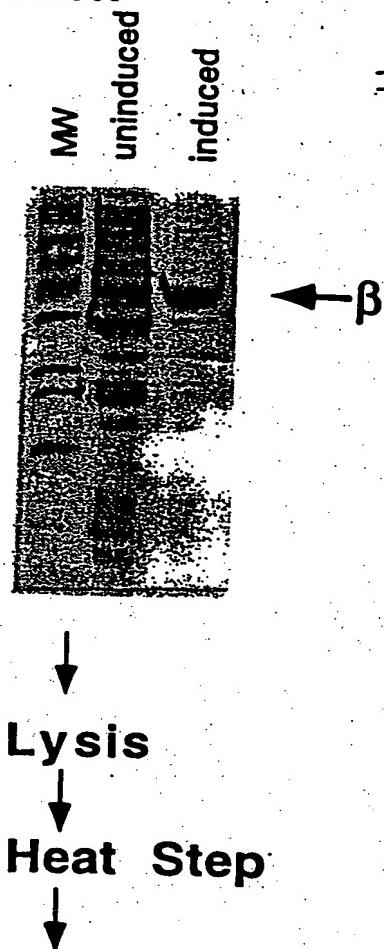


FIG. 24B MonoQ Column

Fraction: 5 7 9 11 13 15 17 19 21 23 25

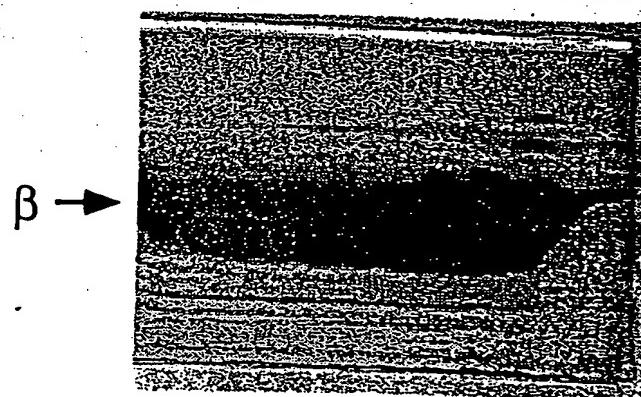
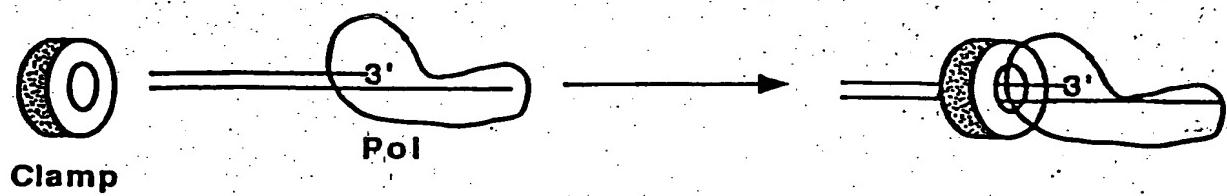


FIG. 25A



Clamp

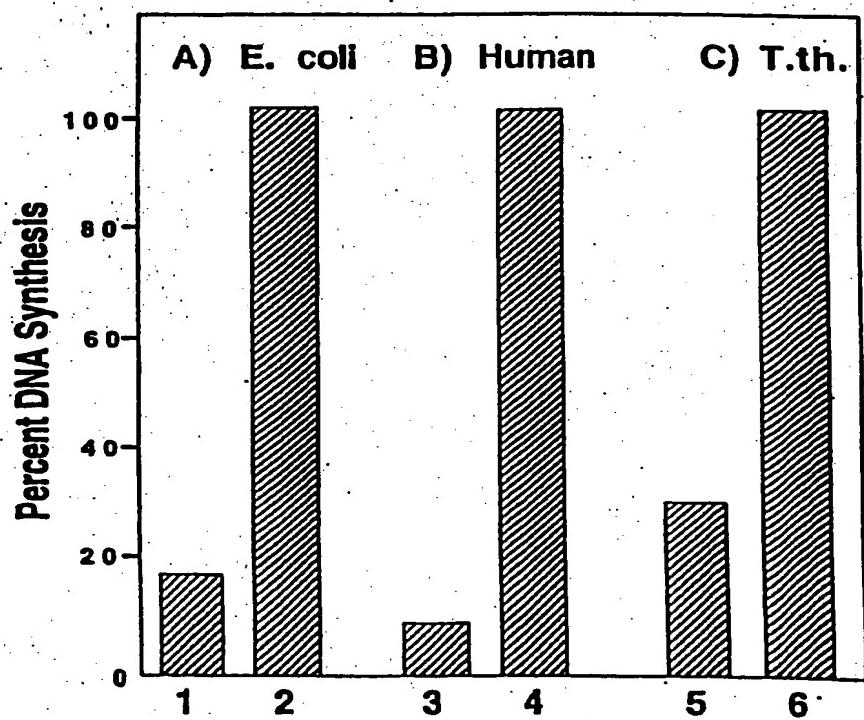


FIG. 25B

FIG. 26A

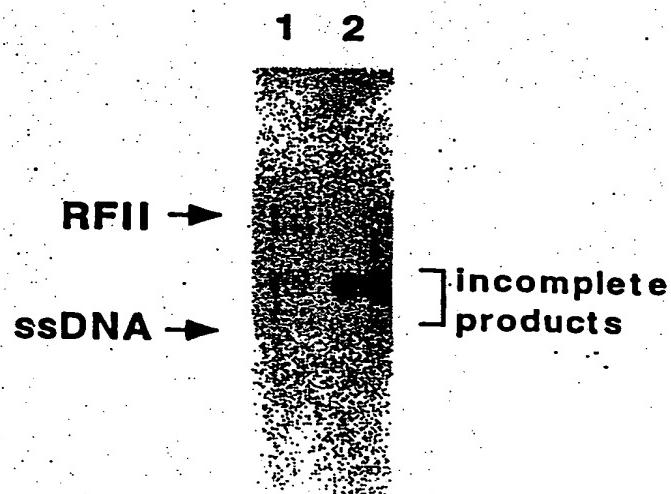
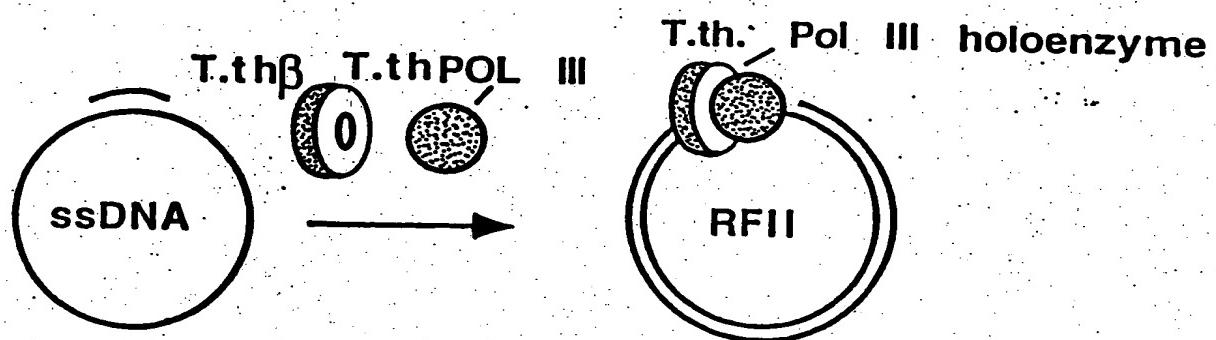


FIG. 26B

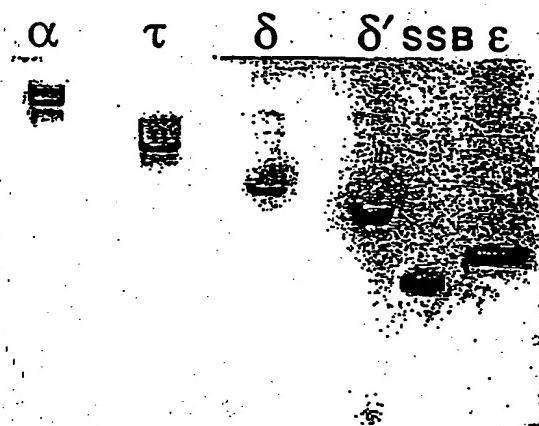


FIG. 27

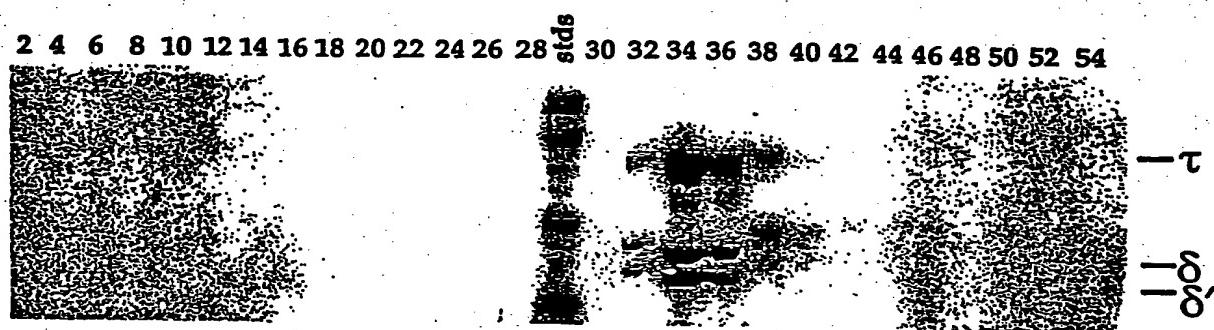


FIG. 28

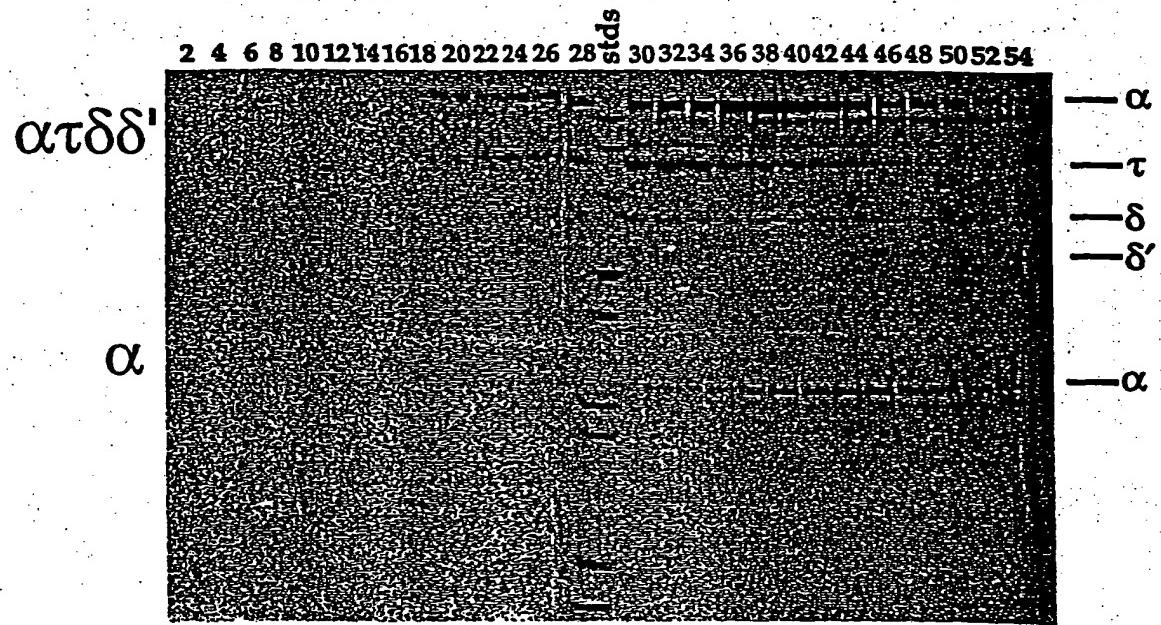


FIG. 29

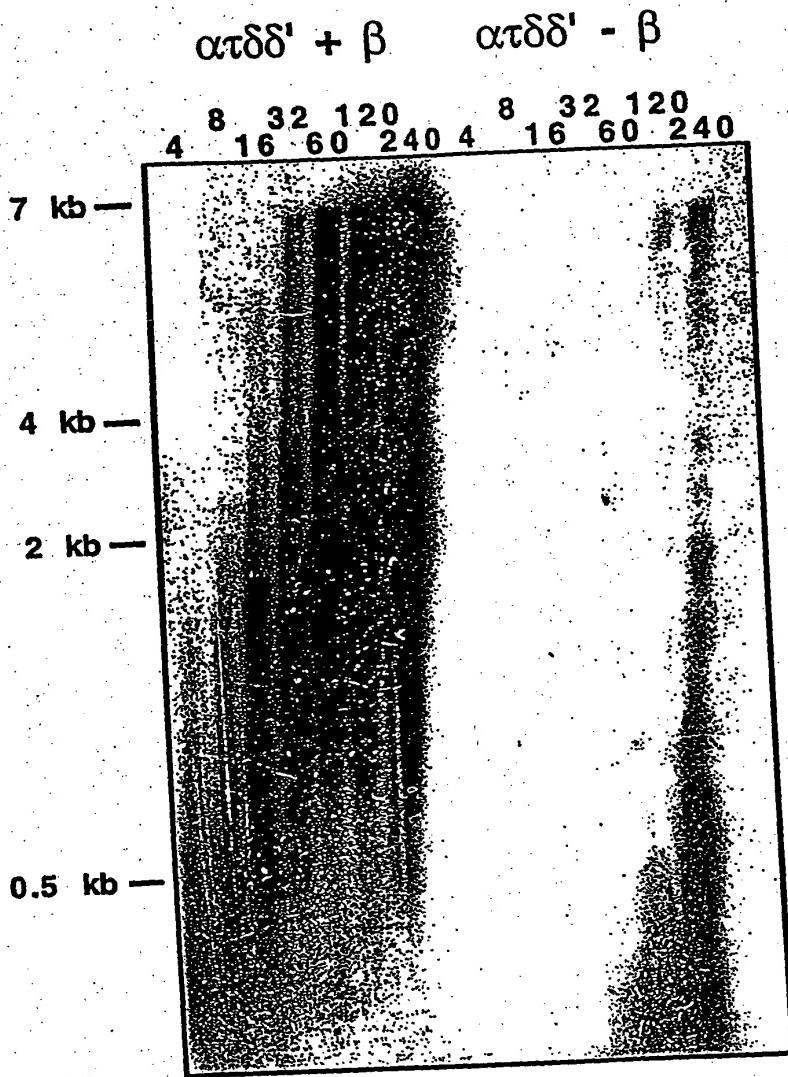


FIG. 30

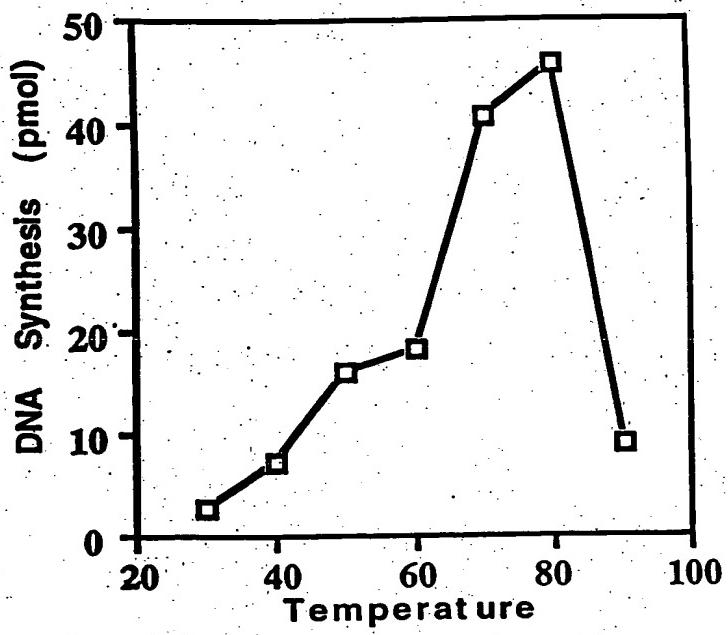


FIG. 31

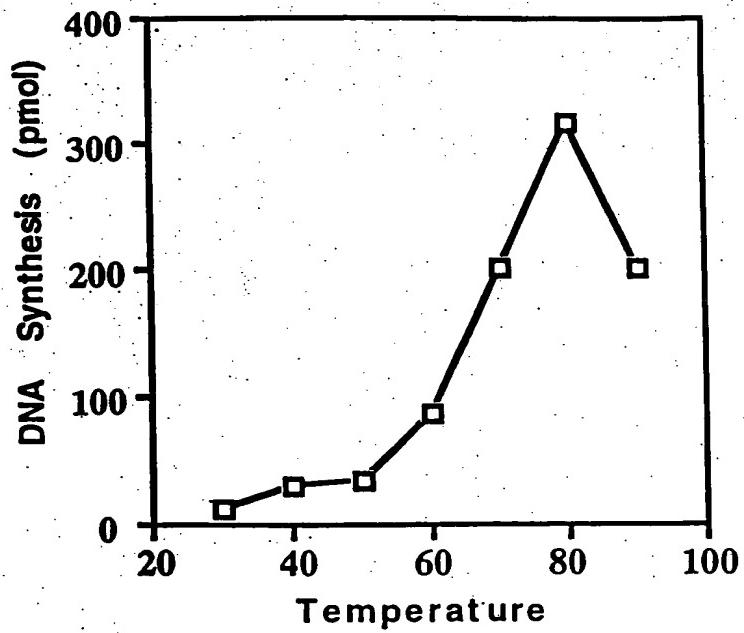
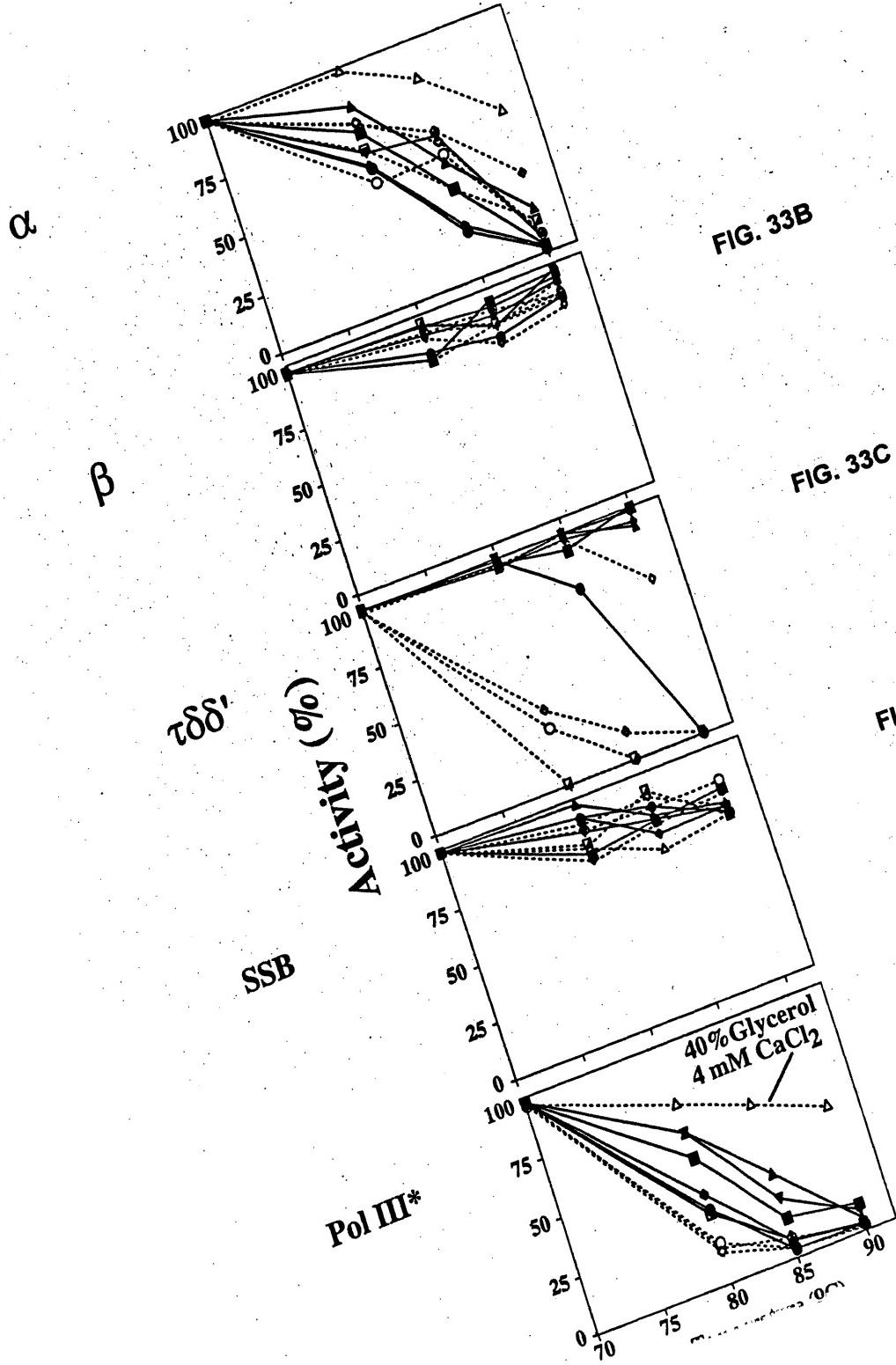


FIG. 32



ATGAGTAAGGATTCTGTCCACCTCACCTGCACACCCAGTTCTCACTCCT	100
GGACGGGGCTATAAAGATAGACGAGCTCGTAAAAAGGCAAAGGAGTATG	
GATACAAAGCTGTGGAATGTCAGACCACGGAAACCTCTTCGGTTCGTAT	200
AAATTCTACAAAGCCCTGAAGGGCGGAAGGAATTAAAGCCCATAATCGGCAT	
GGAAGCCTACTTTACCACGGGTTCGAGGTTGACAGAAAGACTAAAACGA	
GCGAGGACAACATAACCGACAAGTACAACCACCCACTCATACTTATAGCA	300
AAGGACGAAAAGGTCTAAAGAACTTAATGAAGCTCTCAACCCCTCGCCTAC	
AAAGAAGGTTTTACTACAAACCCAGAATTGATTACGAACCTCTTGAAGGAA	400
GTACGGGAGGGCCTAATAGCCCTTACCGCATGCCTGAAAGGTGTTCCCA	
CCTACTACGCTTCTATAAACGAAGTGAAAAAGGCGGAGGAATGGTAAAG	500
AAGTTCAAGGATATATTGGAGATGACCTTATTTAGAACTTCAAGCGAA	
CAACATTCCAGAACAGGAAGTGGCAAACAGGAACAGGAACTTAATAGAGATAGCCA	600
AAAAGTACGATGTGAAACTCATAGCGACGCAGGACGCCACTACCTCAAT	
CCCGAAGACAGGTACGCCACACGGTTCTATGGCACTTCAAAATGAAAAA	700
GACCATTACGAACACTGAGTTGGGAAACTTCAAGTGTCAAACGAAGACC	
TTCACTTTGCTCCACCCGAGTACATGTGGAAAAGTTGAAGGTAAGTTC	800
GAAGGCTGGAAAAGGCACTCCTGAACACTCTCGAGGTAATGGAAAAGAC	
AGCGGACAGCTTGAGATATTTGAAAACCTCACCTACCTCCTTCCCAAGT	900
ACGACGTCCGCCGACAAAACCCCTGAGGAATACCTCAGAGAACTCGCG	
TACAAAGGTTTAAGACAGAGGATAGAAAGGGACAAGCTAAGGATACTAA	1000
AGAGTACTGGGAGAGGCTCGAGTACGAACCTGGAAAGTTATAAAACAAATGG	
GCTTTGGGGATACTTCTTGATAGTTAGGACTTCATAAAACTGGCTAAG	1100
AAAAACGACATACCTGTTGGACCCCGAAGGGGAAGTGTGGAGGTTCCCT	
CGTCGCATACGCCATCGGAATAACGGACGTTGACCTATAAAGCACGGAT	1200
TCCTTTTGAGAGGTTCTAAACCCGAAAGGGTTCCATGCCGGATATA	
GACGTGGATTTCTGTCAGGACAACAGGAAAAGGTATAGAGTACGTAAG	1300
GAACAAAGTACGGACACGACAACGTAGCTCAGATAATCACCTACAACGTA	
TGAAGGCGAACAAACACTGAGAGACGTCGAAGGGCATGGACTCCCC	1400
TACTCCACCGCGGACAAACTCGAAAACCTCATTCCTCAGGGGGACGTTCA	
GGGAACGTGGCTCAGTCTGGAAAGAGATGTACAAACGCCTGTGGAGGAAC	1500
TCCTTCAGAAGTACGGAGAACACAGAACGGACATAGAGGACAACGTAAG	
AAGTTCAAGACAGATATGCGAAGAAAGTCCGGAGATAAAACAGCTCGTTGA	1600
GACGGCCCTGAAGCTTGAAAGGTCTCACGAGACACACCTCCCTCCAGCCG	
CGGGAGTGGTTATGCAACAAAGCCCTTGAGCGAGCTCGTCCCCTCTAC	1700
TACGATAAAAGAGGGCGAAGTCGCAACCCAGTACGACATGGTTCAGCTCGA	
AGAACTCGGTCTCCTGAAGATGGACTTCCTCGGACTCAAAACCCCTCACAG	1800
AACTGAAACTCATGAAAGAAACTCATAAAGGAAAGACACGGAGTGGATATA	
AACTCCCTGAACTTCCCTTGACGACCCGAAAGTTACAAACTCCTTCA	1900
GGAAGGAAAACACGGGAGTGTCCAGCTCGAAAGCAGGGGAATGAAAG	
AACTCCTGAAGAAACTAAAGCCGACAGCTTGACGACATCGTTGGTC	2000
CTCGCACTCTACAGACCCGGACCTCTAAAGAGCGGACTCGTTGACACATA	
CATTAAGAGAAAGCACGGAAAAGAACCCGTTGAGTACCCCTCCCGGAGC	2100
TTGAACCCGTCCTTAAGGAAACCTACGGAGTAATCGTTATCAGGAACAG	
GTGATGAAGATGTCTCAGATACTTCCGGTTACTCCCGGAGAGGGCGGA	2200
TACCCCTCAGAAAGGCAGTAGGTAAGAAGAAAGCGGATTTAATGGCTCAGA	
TGAAAGACAAGTTCATACAGGGAGCGGTGAAAGGGGATACCCTGAAGAA	2300
AAGATAAGGAAGCTCTGGGAAGACATAGAGAAGTTCGCTTCTACTCCTT	
CAACAAGTCTCACTCGGTAGCTTACGGGTACATCTCCTACTGGACCGCCT	2400

FIG. 34A

ACGTTAAAGCCCACATATCCCGGGAGTTCTCGCGGTAAAACTCACA	ACT	
GAAAAGAACGACAACAAGTTCTCAACCTCATAAAAGACGCTAA	ACTCTT	2500
CGGATTGAGATACTTCCCCCGACATAAACAAAGAGTGATGTA	GGGATT	2600
CGATAGAAGGTGAAAACAGGATAAGGTTGGGCTTGGCAGGATAA	AGGGAA	2700
GTGGGAGAGGAAACTGCTAACGATAATCGTTGAAGCTAGAAAGA	ACTATAA	2800
GCAGTTCAAAGGGCTTGGGACTTCATAAACAAAACCAAGAACAG	GAAGAAGA	2900
TAAACAAAGAAAGTCGTGGAAGCACTCGTAAAGGCAGGGCTTGT	ACTTT	3000
ACTAACGAAAGAGGAAAGAAACTACTCGTAAAGTGGCAA	ACTCTGAAAA	3100
AGCATTAAATGGCTACACAAAACCTCCCTTCGGTGCACC	GAAAGAAGAAG	3200
TGGAAGAACACTCGACCCCTTAAAGCTTGAAGAAGTTCTCGG	TTTAC	3300
ATTTCAAGGCACCCCTTGACAACTACGAAAAGCTCCTCAAGA	ACCGCTA	3400
CACACCCATTGAAGATTAGAAGAGTGGGACAAGGAAAGCGAAG	CGCGGTGC	3500
TTACAGGAGTTATCACGGAACCTCAAAGTAAAAAGACGAA	AAACCGGAGAT	
TACATGGCGGTCTCAACCTCGTTGACAAGACGGGACTAATAG	AGTGTGT	
CGTCTTCCGGGAGTTACGAAGAGGCAAAGGAAC	TGATAGAAGAGGACA	
GAGTAGTGGTAGTCAAAGGTTCTGGACGAGGACCTTGAA	ACGGAAAAT	
GTCAAGTTCGTGGTGAAGAGGTTCTCCCTGAGGAGTTCGC	AAAGGA	
GATGAGGAATACCCTTATATATTCTAAAAGAGAGCAAGCC	CTAAACG	
GCCTTGCAGAAAACCTAAAGGAATTATTGAAAACAACAGG	ACGGAGGAC	
GGATACAACCTGGTTCTCACGGTTGATCTGGGAGACTACT	TCGTTGATT	
AGCACTCCCACAAGATATGAAACTAAAGGCTGACAGAAAGG	TGTAGAGG	
AGATAGAAAAACTGGGAGTGAAGGTCTAAATTAGTAAATAAC	CCCTTACT	
TCCGAGTAGTCCCC		

FIG. 34B

MSKDFVHLHLHTQFSLLGAIKIDELVKKAKEYGYKAVGMSDHGNLFGSY	100
KFYKALKKAEGIKPIIIGMEAYFTTGSRFDRKTKTSEDNITDKYNHHLILIA	
KDDKGLKNLMKLSTLAYKEGFYYKPRIDYLELEKYGEGLIALTACLKGP	200
TYYASINEVKKAEWVKKFKDIFGDDLYLELQANNIPEQEVARNLIEIA	
KKYDVKLIATQDAHYLNPEDRYAHTVLMALQMKTIHESGNFKCSNED	300
LHFAPPEMWKKFEGKFEKGWEKALLNTLEVMEKTADSFEIFENSTYLLPK	
YDVPPDKTLEEYLRELAYKGLRQRIERGQAKDTKEYWERLEYELEVINKM	400
GFAGYFLIVQDFINWAKKNDIPVGPGGRGSAGGSLVAYAIGITDVPDIKG	
FLFERFLNPERVSMRDIDVDFCQDNREKVIEYVRNKYGHDNVAQIITYNV	500
MKAQTLRDVARAMGLPYSTADKLAKLIPQGDVQGTWLSLEEMYKTPVEE	
LLQKYGEHRTDIEDNVKKFRQICEESPEIKQLVETALKLEGLTRHTSLHA	600
AGVVIAPKPLSELVPLYDKEGEVATQYDMVQLEELGLLKMDFLGLKTLT	
ELKLMKELIKERHGVDFINFELPLDDPKVYKLLQEGKTTGVFQLESRGMK	700
ELLKKLKPDSDFDDIVAVLALYRPGPLKSGLVDTYIKRKHGKEPVEYPFPE	
LEPVLKETYGVIVYQEVMKMSQILSGFTPGEADTLRKAIGKKADLMAQ	800
MKDQFIQGAVERGYPEEKIRKLWEDIKFASYSFNKSHSVAYGYISWT	
YVKAHYPAAFFAVKLTTEKNDNKFLNLIKDAKLFGEIELPPDINKSDVGF	900
TIEGENRIRFGLARARIKGVEETAKIIVEARKKYQFKGLADFINTKNRK	
INKKVVEALVKAGAFDFTKKRKELLAKVANSEKALMATQNSLFGAPKEE	1000
VEELDPLKLEKEVLGFYISGHPLDNYEKLLKNRYTPIEDLEEWDKES	
LTGVITELKVKKTKNGDYMMAVFNLVDKTGLIECVVFPGVYEAKELIEED	1100
RVVVKGFLDEDLETENVKFVVKEVFSPEEFAKEMRNTLYIFLKREQALN	
GVAEKLKGIIENNRTEDGYNLVTVLDLGDFVDLALPQDMKLKADRKVVE	
EIEKLGVKVII	1161

FIG. 35

ATGAACTACGTTCCCTCGCGAGAAAGTACAGACCGAAATTCTTCAGGGAA	100
AGTAATAGGACAGGAAGCTCCCGTAAGGATACTCAAAAACGCTATAAAAAA	
ACGACAGAGTGGCTCACGCCAACCTCTTGCCGGACCGAGGGGGTTGGGAA	200
AAGACGACTATTGCAAGAATTCTCGAAAAGCTTGAACGTAAAAATCCCTC	
CTCCAAAGGTGAGCCCTGCGGTGAGTGCAGAAAACGAGATAGACA	300
GGGGTGTGTTCCCTGACTTAATTGAAATGGATGCCCTCAAACAGGGT	
ATAGACGACGTAAGGCATTAAAAGAAGCAGGTCAATTACAAACCTATAAA	400
AGGAAAGTACAAGGTTACATAATAGACGAAGCTCACATGCTACGAAAG	
AAGCTTCAACGCTCTCTTAAACCCCTCGAAGAGCCCCCTCCCAGAACT	500
GTTTCGTCTTGTACACGGAGTACGACAAAATTCTCCCACGATACT	
CTCAAGGTGTCAGAGGATAATCTCTCAAAGGTAAAGAAAGGAAAAGTAA	600
TAGAGTATCTAAAAAGATATGTGAAAAGGAAGGGATTGAGTGCAGAG	
GGAGCCCTTGAGGTTCTGGCTCATGCCCTCTGAAGGGTGCATAGGGATGC	700
AGCCTCTCTCCCTGGACCAGGCAGCGTTACGGGGAGGCAGGGTAACAA	
AAGAAGTAGTGGAGAACTTCCTCGAATTCTCAGTCAGGAAAGCGTTAGG	800
AGTTTCTGAAATTGCTTCTGAACCTCAGAAGTGGACGAAGCTATAAGTT	
CCTCAGAGAACTCTCAGAAAAGGGTACAAACCTGACCAAGTTGGAGA	900
TGTTAGAAGAGGAAGTGAGAAACGCAATTAGTAGTAAAGAGCCTGAAAAT	
CCCGAAAGCGTGGTTAGAAGTGGCAGGATTACGAAGACTTCAAAGACTA	1000
CCCTCTGGAAGCCCTCTACGTTGAGAACCTGATAAACAGGGTAAAG	
TTGAAGCGAGAACGAGAGAACCTTAAGAGCCTTGAACCTCGCGTAATA	1100
AAGAGCCTTATAGTCAAAGACATAATTCCCGTATCCAGCTCGGAAGTGT	
GGTAAAGGAAACCAAAAGGAAGAAAAGAAAGTTGAAGTAAAAGAAGAGC	1200
CAAAGTAAAAGAAGAAAACCAAAAGGAGCAGGAAGAGGACAGGTTCCAG	
AAAGTTTAAACGCTGTGGACGGAAAATCCTAAAGAATACTTGAAGG	1300
GGCAAAAAGGAAGAAAGAGACGGAAAATCGCTAAAGATAGAACGCT	
CTTATCTGAGAACCATGAAAAGGAATTGACTCACTAAAGGAGACTTT	1400
CCTTTTTAGAGTTGAACCGTGGAGGATAAAAAACCTCAGAACGTC	
CAGCGGGACGAGGCTGTTAAAGGTAAAGGAGCTCTCAATGCAAAAT	1500
ACTCAAAGTACGAAGTAAAGCTAAGGTACATAAGGTGAGAATGCCGTG	
GAAGAGATAGGGCTGTTAACGCACTAATAGACGGCTGCCAGGTACGC	1600
ACTCACGAGGACGAAGGAAAAGGGAAAGGGAGAAGTTTCGTTTAGCGA	
CTCCTTATAAAAGTCAAGGAATTGATGGAAGCTATGGAGGGTATGAAAAAA	1700
CACATAAAAGGATTAGAAATCTCGGAGAGACGGATGAGGATTAACTTT	
TTAAAGTATGGGTGTATCTGAGCAAAGGTTAACGCTAAACAAACCTGA	1800
AACCCGCAAGGGACCGCCGAAAGCCATAAAAAACTCCTGAAAAACCTA	
AGGAAAGGCATAAAAGAACAAACACTCTCGGAGTCACGGGAAGCGGAAA	1900
GACTTTACTCTAGCAAACGTAATAGCGAAGTACAACAAACCAACTCTG	
TGGTAGTTACAAACAAAATTCTCGCGCACAGCTATACAGGGAGTTAAA	2000
GAACATTCCTGAAAACGCTGTAGAGTACTTGTCTCTTACTACGACTA	
TTACCAACCTGAAGCCTACATTCCGAAAAAGATTATACATAGAAAAGG	2100
ACGCGAGTATAAACGAAAGCTGGAACGTTCAAGACACTCCGCCACGATAT	
CCGTTCTAGAAAGGAGGGACGTTAGTGTAGTGTCTCAGTTCTGCATA	2200
TACGGACTCGGGAAACCTGAGCACTACGAAAACCTGAGGATAAAACTCCA	
AAGGGGAATAAGACTGAACCTGAGTAAGCTCCTGAGGAAACTCGTTGAGC	2300
TAGGATATCAGAGAAATGACTTGCCTAAAGAGGGCTACCTTCTCGGTT	
AGGGGAGACGTGGTTGAGATAGTCCCTCTCACACGGAAGATTACCTCGT	2400
GAGGGTAGAGTTCTGGGACGACGAAGTTGAAAGAATAGTCCTCATGGACG	
CTCTGAAC	

FIG. 36

MINYVPFARKYRPKFFREVIGQEAPVRILKNAIKNDRVAHAYLFAGPRGVG	
KTTIARILAKALNCKNPSKGEPCECENCREDRGVFPDLIEMDAASNRG	100
IDDVRALKEAVNYKPIKGKYKVYIDEAHMLTKEAFNALLKTLEEPPRT	
VFVLCTTEYDKILPTILSRCQRIIFSKVRKEVKIEYLKKICEKEGIECEE	200
GALEVLAHASEGCMRDAASLLDQASVYGEGRVTKEVENFLGILSQESVR	
SFLKLLLNLSEVDEAIKFRLRELSEKGYNLTFWEMLEEVRNAILVSLKN	300
PESVVQNWDYEDFKDYPLEALLYVENLINRGKVEARTREPLRAFELAVI	
KSLIVKDIIIPVSQSQLGSVVKETKKEEKVVEKEEPKVKEEKPKEQEEDRFQ	400
KVLAvgkilkRilegakreerDGKIVLkieasylrtmkkefdslketeF	
PFLEFEPVEDKKPKQKSSGTRLF	473

FIG. 37

ATGCGCGTTAAGGTGGACAGGGAGGAGCTTGAAGAGGTTCTTAAAAAAGC	100
AAGAGAAAAGCACGGAAAAAAAGCCGCACTCCCATACTCGCGAACTTCT	
TACTCTCCGCAAAAGAGGAAAACCTTAATCGTAAGGGCAACGGACTTGGAA	200
AACTACCTTGTAGTCTCCGTAAGGGGAGGTTGAAGAGGAAGGAGAGGT	
TTGCGTCCACTCTCAAAACTCTACGATATAGTCAGAACCTAAATTCCG	
CTTACGTTACCTTCATACGGAAGGTGAAAAACTCGTCATAACGGGAGGA	300
AAGAGTACGTACAAACTTCCGACAGCTCCCGCGGAGGACTTCCGAATT	
TCCAGAAATCGTAGAAGGAGGAGAACACTTTCGGAAACCTTCTCGTTA	400
ACGGAATAGAAAAGGTAGAGTACGCCATAGCGAAGGAAGCGAACATA	
GCCCTTCAGGGATGTATCTGAGAGGATACGAGGACAGAATTCACTTTGT	500
GTTCCGGACGGTCACAGGCTTGCACCTTATGAACCTCTACGTAACATTGA	
AAAGAGTGAAGACGAGTCCTTGCTTACTTCTCCACTCCGAGTGGAAAC	600
TCGCCGTTAGCTCCTGGAAGGAGAATTCCCGACTACATGAGTGTCATCC	
CTGAGGAGTTTCGGCGGAAGTCTTGTGAGACAGAGGAAGTCTTAAAG	700
GTTTTAAAGAGGTTGAAGGCTTAAGCGAAGGAAAAGTTTCCCGTGAA	
GATTACCTTAAGCGAAAACCTTGCATCTTGAGTCGCGGATCCGGAGT	800
TCGGAGAAGCGAGAGAGGAAATTGAAGTGGAGTACACGGGAGAGCCCTT	
GAGATAGGATTCAACGAAATACCTTATGGAGGCGCTTGACGCCCTACGAC	900
AGCGAAAAGAGTGTGGTTCAAGTTACAACCCCCGACACGCCACTTATT	
GGAGGCTGAAGATTACGAAAAGGAACCTTACAAGTGCATAATAATGCCGA	1000
TGAGGGTGTAGCCATGAAAAAGCTTTAATCTTTTATTGAGCTTGAGCC	
TTTTAATTCCCTGCGTTAGCGAAGCCAAACCCAAAGTCTTC	1090

FIG. 38

MRVKVDREEEVLKKARESTEKKAAALPILANFLSAKEENLIVRATDLE	100
NYLVVSVKGEVEEEGEVCVHSQKLIDIVKNLNSAYVYLHTEGEKLVITGG	
KSTYKLPTAPAEDFPEFPEIVEGETLSGNLLVNGIEKVEYAIKEEANI	200
ALQGMYLRGYEDRIHFVGSDGHRLALYEPLGEFSKELLIPRSLKVLKKL	
ITGIEDVNIEKSEDESFAYFSTPEWKLAVRLLEGEPDYSMSVIPEEFSAE	300
VLFETEEVLKVLKRLKALSEGKVFPVKITLSENLAIFEFADPEFGEAREE	
IEVEYTGEFIEGFNGKYLMEALDAYDSERVWFKFTTPDTATLLEAEDYE	
KEPYKCIIMPMRV	363

FIG. 39

GTGGAAACCACAATATTCCAGTTCAGAAA	100
GAAGGAGAGGGTCTCGTCCTCATGGAGAAGAGCAGTATCTCATAAGAA	
CCTTTTGTCTAAGCTAAGGAAAAGTACGGGGAGAATTACACGGTTCTG	200
TGGGGGGATGAGATAAGCGAGGAGGAATTCTACACTGCCCTTCCGAGAC	
CAGTATATTCGCGGTTCAAAGGAAAAGCGGTGGTCATTACAACCTCG	300
GGGATTCCCTGAAGAACGCTCGGAAGGAAGAAAAAGGAAAAGAAAGGCTT	
ATAAAAGTCCTCAGAACGTAAAGAGTAACACTACGTATTATAGTGTACGA	400
TGCGAAACTCCAGAACAGGAACCTTCAGGAACCTCTGAAATCCGTAG	
CGTCTTCGGCGGTATAGTGGTAGCAAACAGGCTGAGCAAGGAGAGGATA	500
AAACAGCTCGTCCTTAAGAAGTTCAAAGAAAAGGGATAAACGTAGAAA	
CGATGCCCTTGAAATACCTTCTCCAGCTCACGGTTACAACCTGATGGAGC	600
TCAAACATTGAGGTTGAAAAACTGATAGATTACGCAAGTGAAAAGAAAATT	
TTAACACTCGATGAGGTAAAGAGAGTAGCCTCTCAGTCTCAGAAAACGT	700
AAACGTATTGAGGTTCGTTGATTACTCCTCTTAAAGATTACGAAAAGG	
CTCTTAAAGTTTGGACTCCCTCATTCCTCGGAATACACCCCCCTCCAG	800
ATTATGAAAATCCTGTCCCTCATGCTCTAAACTTACACCCCTCAAGAG	
GCTTGAAGAGAAGGGAGAGGACCTGAATAAGGCATGGAAAGCGTGGGAA	900
TAAAGAACAACTTCTCAAGATGAAGTTCAAATCTTACTTAAAGGCAAAC	
TCTAAAGAGGACTTGAAGAACCTAATCCTCTCCCTCAGAGGATAGACGC	1000
TTTTCTAAACTTACTTCAGGACACAGTCAGTTGCTGGGATTCTT	
GACCTCAAGACTGGAGAGGAAAGTTGTGAAAATACTCTCATGGTGGAT	
AATCTTTTATGAAGTTGCAGTTGCCTTCCCAGTTCT	1093

FIG. 40

VETTIQFQKTFTKPPKERVFVLHGEEQYLIRTFLSKLKEKYGENYTVL	100
WGDEISEEEFYTALSETSIFGGSKEKAVVIYNFGDFLKKLGRKKKEKERL	
IKVLRNVKSNYVFIVYDAKLQKQELSSEPLKSVASF GGIVVANRLSKERI	200
KQLVLKKFKEKGINVENDALEYLLQLTGYNLMELKLEVEKLIDYASEKKI	
LTLDEVKRVAFSVSENVNFVFVDLLLKDYEKALKVLDLSLISFGIHPLQ	300
IMKILSSYALKLYTLKRLEEKGEDLNKAMESVGIKNNFLKMKFKSYLKAN	
SKEDLKNLILSLQRIDAFLSKLYFQDTVQLLRFITSRLEREVVKNTSHGG	

FIG. 41

ATGGAAAAAGTTTTTGAAAAACTCCAGAAAACCTGCACATACCCGG
 AGGACTCCTTTACGGCAAAGAAGGAAGCGGAAAGACGAAAACAGCTT
 TTGAATTGCAAAAGGTATTTATGTAAGGAAAACGTACCTGGGATGCG
 GAAGTTGTCCCTCTGCAAACACGTAAACGAGCTGGAGGAAGCCTTCTT
 AAAGGAGAAATAGAAGACTTTAAAGTTATAAGACAAGGACGGTAAAAG
 CACTTCGTTTACCTTATGGCGAACATCCCACCTTGTGGTAATAATCCC
 GAGCGGACATTACATAAAGATAGAACAGATAAGGAAAGTTAAGAACTTTG
 CCTATGTGAAGCCCGACTAAGCAGGAGAAAAGTAATTATAATAGACGAC
 GCCCACGCGATGACCTCTCAGGCGCAAACGCTCTTTAAAGGTATTGGA
 AGAGCCACCTGCGGACACCACCTTATCTTGACCACGAACAGGCCTCTG
 CAATCCTGCCGACTATCCTCTCAGAACCTTCAAGTGGAGTTCAAGGGC
 TTTTCAGTAAAAGAGGTTATGAAATAGCGAAAGTAGACGAGGAAATAGC
 GAAACTCTCTGGAGGCAGTCTAAAAGGGCTATCTTACTAAAGGAAAACA
 AAGATATCCTAAACAAAGTAAAGGAATTCTGGAAAACGAGCCGTTAAA
 GTTTACAAGCTTGCAGTGAATTGAAAAGTGGGAACTGAAAAGCAAAA
 ACTCTTCTTGAATTATGGAAGAATTGGTATCTCAAAATTGACCGAAG
 AGAAAAAAAGACAATTACACCTACCTTCTTGATACGATCAGACTCTTAAA
 GACGGACTCGCAAGGGGTGAAACGAACCTCTGTGGCTTACGTTAGC
 CGTTCAGGCGGATTAATAAAACGTTATTGATTCCGTAACATTAAACCTT
 AATCTAAATTATGAGAGCCTTGAAGGAGGTCTGGTATGAAATTGAA
 GATTAGATATATAGATACGAGGAAGATAGGAACCGTGAGCGGTGTAAAAG
 T

FIG. 42

MEKVFLEKLQKTLHIPGGLLFYKGEGSGKTAFEFAKGILCKENVPWGC
 GSCPSCHVNELEEAFFKGEIEDFKVYKDKGKKHFVYLMGEHPDFVVII
 PSGHYIKIEQIREVKNFAYVKPALSRRKVIIDDAHAMTSQAANALLKVL
 EEP PADTFILTTNRRSAILPTILSRTFQVEFKGSVKEVMEIAKVDEEI
 AKLSGGSLKRAILLKENKDILNKVKEFLENEPLKVYKLASEFEKWEPEKQ
 KLFLEIMEELVSQKLTEEKKDNYTYLLDTIRLFKDGLARGVNEPLWLFTL
 AVQAD

FIG. 43

ATGAACTCCTGAAAAGTCCTTTACTGAGAAAAGCTAAAAGTCTCC	
TTACTTCGAAGAGTTCTACGAAGAAATCGATTGAAACCAGAAGGTGAAAG	100
ATGCAAGGTTGTAGTTTGACTGCGAACAGAACTCGACGTAAAG	
AAGGCAAAACTCCTTCAATAGGTGCGGTTGAGGTTAAAACCTGGAAAT	200
AGACCTCTCTAAATCTTTACGAGATACTCAAAGTGACGAGATAAAGG	
CGGCGGAGAGATAATGGAATAACCAGGGAAAGACGTTGAAAAGTACGGAAAG	300
GAACCAAAGGAAGTAATATACGACTTCTGAAGTACATAAAGGGAAAGCGT	
TCTCGTTGGCTACTACGTGAAGTTGACGTCTCACTCGTTGAGAAGTACT	400
CCATAAAAGTACTTCCAGTATCCAATCATCAACTACAAGTTAGACCTGTT	
AGTTTGTGAAGAGAGAGTACCAAGTGGCAGGAGTCTGACGACCTTAT	500
GAAGGAACTCGGTGTAGAAATAAGGGCAAGGCACAACGCCCTGAAGATG	
CCTACATAACCGCTCTCTTCTAAAGTACGTTACCGAACAGGGAG	
TACAGACTAAAGGATCTCCGATTTCCTT	600

FIG. 44

MNFLKKFLLLRKAQKSPYFEFYEEIDLNLQKVNDARFVVFDCEATELDVK	
KAKLLSIGAVEVKNLEIDLSSKFYEILKSDEIKAAEIHGITREDVEKYGK	100
EPKEVIYDFLKYIKGSVLVGYVFKFDVSLVEKYSIKYFQYPIINYKLDLF	
SFKREYQSGRSIQLLMKELGVEIRARHNALEDAYITALLFLKYVYPNRE	200
YRLKDLPIFL	

FIG. 45

ATGCTCAATAAGGTTTTATAATAGGAAGACTTACGGGTGACCCCGTTAT
AACTTATCTACCGAGCGGAACGCCGTAGTAGAGTTACTCTGGCTTACA 100
ACAGAAGGTATAAAAACCAGAACGGTGAATTTCAGGAGGAAAGTCACCTC
TTGACGTAAGAGCGTACGGAAAAATGGCTGAAGACTGGGCTACACGCTT
CTCGAAAGGATACCTCGTACTCGTAGAGGGAAGACTCTCCCAGGAAAAGT 200
GGGAGAAAGAAGGAAAGAAGTTCTCAAAGGTCAAGGATAATAGCGGAAAAC
GTAAGATTAATAAACAGGCCGAAAGGTGCTGAACCTCAAGCAGAAGAAGA 300
GGAGGAAGTTCTCCCATTGAGGAGGAAATTGAAAAACTCGGTAAAGAGG
AAGAGAAGCCTTTACCGATGAAGAGGGACGAAATACCTTTAATTGTAGA 400
GGAGGTTAAAGTATGGTAGTGAGAGCTCTAAGAAGAAAGTTGTATGTA
CTGTGAACAAAAGAGAGAGCCAGATT 500

FIG. 46

MLNKVFIIGRLTGDPVITYLPSGTPVVEFTLAYNRRYKNQNGEFQEESHF
FDVKAYGKMAEDWATRFSKGYLVLVEGRLSQEKWEKEGKKFSKVRIIAEN 100
VRLINRPKGAELOAEEEEEVPPPIEEEIEKLGKEEEKPFTDEEDEIPF

FIG. 47

ATGCAATTGTGGATAAACTTCCCTGTGACGAATCCGCCAGAGGGCGGT	100
TCTTGGCAGTATGCTTGAAGACCCGAAAACATACTCTGGTACTTGAAT	
ACCTTAAAGAAGAAGACTCTGCATAGACGAGCACAAGCTACTTTCAAGG	200
GTTCTTACAAACCTCTGGTCCGAGTACGGCAATAAGCTCGATTCGTATT	
AATAAAAGGATCACCTTGAAAAGAAAAACTTACTCCAGAAAATACCTATAG	
ACTGGCTCGAAGAACTCTACGAGGAGGCGGTATCCCCCTGACACGCTTGAG	300
GAAGTCTGCAAATAGTAAAACAACGTTCCGCACAGAGGGCGATAATTCA	
ACTCGGTATAGAACTCATTCACAAAGGAAAGGAAAACAAAGACTTTCACÀ	400
CATTAATCGAGGAAGGCCAGAGCAGGATATTTCCATAGCGGAAAGTGCT	
ACATCTACGCAGTTTACCATGTGAAAGACGTTGCGGAAGAAGTTATAGA	500
ACTCATTATAAATTCAAAGCTCTGACAGGCTAGTCACGGACTCCCAA	
GC GGTTTCACGGAAC TCGATCTAACAGACGACGGGATTCCACCTGGAGAC	600
TTAATAATACTCGCCGCAAGACCCGGTATGGGAAAACCGCCTTATGCT	
CTCCATAATCTACAATCTCGCAAAGACGAGGGAAAACCTCAGCTGTAT	700
TTTCCTGGAAATGAGCAAGGAACAGCTCGTTATGAGACTCCTCTCTATG	
ATGTCGGAGGTCCCACTTCAAGATAAGGTCTGGAAGTATACTGAATGA	800
AGATTTAAAGAAGCTTGAAGCAAGCGCAATAGAACTCGCAAAGTACGACA	
TATACCTCGACGACACACCCGCTCTCACTACAACGGATTAAAGGATAAGG	900
GCAAGAAAGCTCAGAAAGGAAAAGGAAGTTGAGTTCGTGGCGGTGGACTA	
CTTGCAACTCTGAGACGCCAGTCCGAAAGAGTTCAAGACAGGAGGAAG	1000
TGGCAGAGGTTCAAGAAACTTAAAGCCCTGCAAAGGAACCTCACATT	
CCCGTTATGGCACTTGCAGCTCTCCGTGAGGTGGAAAAGAGGAGTGA	1100
AAAAGACCCAGCTTGCAGCTCAGAGAATCCGACAGATAGAACAGG	
ACGCAGACCTAACCTTTCCACAGACCCGAGTACTACAAGAAAAAG	1200
CCAAATCCCGAAGAGCAGGGTATAGCGGAAGTGATAATAGCCAAGCAAAG	
GCAAGGACCCACGGACATTGTGAAGCTCGCATTATTAAAGGAGTACACTA	1300
AGTTTGCAAACCTAGAACGCCCTCTGAACAACTCCTGAAGAACAGGAA	
CTTCCGAAATTATTGAACACAGGAGGATGAAGGATTGAAGATATTGA	1400
CTTCTGAAAATTAAAGGTTTATAATTATCTTGGCTATCCGGGTAGCT	
CAATCGGCAGAGCGGGTGGCTG	1472

FIG. 48

MQFVDKLPDES AERAVL GSM LED PEN I PLV LEYL KEED FC IDE HKLL FR	100
VLTNLWSEYGNKLDFVLI KDHLEKKNLLQKIPIDWLEELYEEAVSPDTLE	
EVCKIVKQRSAQR AIIQLGITSTQFYHVKDVAEEVIELIYKFKSSDRLVT	200
GLPSGFTELDLKTTGFHPGDLIILAARPGMGKTA FMLSIIYNLAKDEGKP	
SAVFSLEMSKEQLVMRLLSMMSEVPLFKIRSGSISNEDLKKLEASAIELA	
KYDIYLD DTPALTTD LIRARKLRKEKEVEFVAVDYLQLLRPPVRKSSR	300
QEEVAEVSRNLKALAKELHIPVMA LAQLSREVEKRSDKRPQLADLRESQ	
I EQDADLILFLHRPEYYKKPNPEEQGIAEVIIAKQRQGPTDIVKLAFIK	400
EYTKFANLEALPEQPPEEEELSEIIEQEDEGFEDIDF	

FIG. 49

ATGTCCTCGGACATAGACGA	ACTTAGACGGAA	ATAGATATAG	AGACGT	100	
CATTTCGA	AATACTTAA	ACTTAGAGA	AGGTAGGTTCC	AAATTACAGAACGA	100
ACTGTC	CCCTTCA	CCCTGACG	ATACACCCTC	TTTACGTGTCTCCA	100
AAACAA	ATATTCA	AGTGTTCGGT	TGCGGGTAGGGGAGAC	GCGATAAAA	200
GTTCGT	TTCCCTTAC	GAGGACATCTC	CTATTGAA	AGGCCGCCTTGAA	200
TCGAAA	ACGCTACGGAA	AGAAATTAGAC	CTTGAAAAGA	TATCAAAAGAC	300
GAAAAGGT	TACGTGGCT	TTGACAGGGTTGT	GATTTCTACAGGGAA	AG	300
CCTTCT	AAAAACAGAGAGG	CAAGTGAGTACG	TAAGAGTAGGGGA	ATAG	400
ACCC	TAAGTAGCGAGGA	AGTTGATCTGGT	ACGCACCTC	CAGTGAA	400
GCAC	TCGTAAAAG	TCTAAAAGAGAACG	ATCTTTAGAGG	CTTACAC	500
AACT	AAAACCTC	TTCTACGAAGGGT	TTTACAGGGAT	CTCTTCA	500
TTCGGCGT	GTCGTGATCCC	GATAAAGGATCC	GAGGGGAAGAG	TATAGGT	600
TTCGGT	GGAAAGGAGG	ATAGTAGAGG	ACAATCTCCA	AGTACATAAA	600
TCCAGACAGCAGGGT	TATTAAAAGGGGAGA	ACTTACCGGT	TTACAGG	GGAAAGG	700
AGGCAA	AGGAGTATATAAAGGA	AGAAGGATTGCG	ATACTTGTG	GAAGGG	700
TACTTTGAC	CTTTGAGACTTTT	CCGAGGGATA	AAAGGAAC	GTTGTTGC	800
ACCC	CTCGGTACAGCC	CTGACCCAAA	ATCAGGCAA	ACCTC	800
TCAC	AAAAAAGGTCTAC	ATCCTTACGACGGAGA	TGATGC	GGGAAGAAAG	900
GCT	ATGAAAAGTGC	CATTCCC	TACTCCTAGTGCA	GGAGTGGAA	900
TCCC	CGTTACCTCCCC	GAAGGATACGATCCC	GACGAGTT	TATAAAGGA	1000
TCGGG	AAAGAGGA	TTAAGAAGACTG	ATAAACAGCTC	AGGGAGCT	1000
AAAACG	GCTCATAAAACCG	CAAGGGAAA	ACTTAGAGGAGA	AAACGCGTGA	1100
GTT	CAGGTATTATCTGG	CTTATTTCCGATGG	AGTAAGGC	GCTTGC	1100
TGG	CTCGGAGTT	CACACCAAGT	ACAAAGT	CCCTATGGAA	1200
ATG	AAAATTGAAAAAA	ATTCTCA	AGAAAAGAA	ATAAACTCTC	1200
GG	AAAATCTCCTG	AAAGGACTG	ATAGAATT	AAAACCAAA	1300
TTG	GAAGTCCTGAA	ACTTAAGTC	CTGAGTTAAGGAA	ACTCGCAGTTAACGCC	1300
TTAA	ACGGAGAGGAGC	ATTACTTCC	AAAAGAAGT	TCTCGAGTACCA	1400
GG	ATAACTTGGAGAA	ACTTTAACAA	CATCCTAGGGATT	ACAAAAAT	1400
CTGGG	AAAAGAGGA	AGAAAAGAGGG	TTGAAAATG	TAATAACTTAA	1500
ACTTTA	ATAAAATT	TTAGAGT	TAGGA		1500

FIG. 50

MSSDIDE	LRR	ELRREIDIVDV	ISEYLN	LEKVG	SNYRTNC	CPFH	PDDTPSFY	VSPSKQ	I	100		
I	FKCF	FGCGV	GGDAIKF	VSLYED	ISYFE	AALELA	KRYG	KLD	LEKISKD	100		
EK	V	VALDR	VCDFY	RESL	LN	REASEY	VKS	RGIDPK	VARKFDLG	YAPSSE	200	
A	LV	VKL	KENDL	LEAY	LET	KNLLS	PTKG	VYRDL	FLRR	VIPIKDPRGRVIG	200	
F	GG	RRI	VED	KSP	KYIN	SPDSRV	FKK	GENL	FGLYE	AEKEYIKEEGFAILVEG	300	
Y	F	DLL	RLI	FSEG	IRNVV	APLGTAL	TQNQAN	LSKF	TKKVYILY	DGDDAGRK	300	
A	M	KS	AIPL	LLS	SAG	VEV	PVLP	PEGYDP	DEFIKE	FGKEELR	LISSGELF	400
E	T	LIK	TAREN	LEEK	TREF	FRYYLG	FISDG	VRR	FALASE	HFTKY	KVPMEILL	400
L	K	I	KNSQE	KE	I	LSF	KE	I	FLKGL	IEL	KPKIDLEVNL	498
N	G	E	H	LLP	KEV	LEYQVDN	LEKLFNN	ILRDLQ	KSGKKR	GLKN	VNT	498

FIG. 51

ATGCAAGATAACCGCTACCTGCAGTATTGTCAGGGGACGGGATTGTAAA	100
GACCGAAGACAACAAGGTAAGGCTCTGCAATGCAGGTTCAAGAAAAGGG	
ATGTAACACAGGAACTAAACATCCCAAAGAGGTACTGGAACGCCAACCTA	200
GACACTTACCCACCCAAAGAACGTATCCCAGAACAGGGCACTTTGACGAT	
AAGGGTCTTCGTCCACAACCTCAATCCCAGGAAGGGAAAGGGCTTACCT	300
TTGTAGGATCTCCTGGAGTCGGCAAAACTCACCTTGGGTTGCAACATTA	
AAAGCGATTTATGAGAAGAAGGGAACTCAGAGGATACTTCTCGATAACGAA	400
GGATCTAATATTCAAGGTTAAAACACTTAATGGACGAGGGAAAGGATAACAA	
AGTTTTAAAAACTGTCTAAACTCACCGGTTTGGGTTCTGACGACCTC	500
GGTTCTGAGAGGCTCAGTGAUTGGCAGAGGGAACTCATCTTACATAAT	
CACTTACAGGTATAACAACCTTAAGAGCACGATAATAACCACGAATTACT	600
CACTCCAGAGGGAAAGAAGAGTAGCGTGAGGATAAGTGCAGGATCTGCA	
AGCAGACTCGGAGAAAACGTAGTTCAAAAATTACGAGATGAACGAGTT	700
GCTCGTTATAAAGGGTCCGACCTCAGGAAGTCTAAAAGCTATCAACCC	
CATCT	

FIG. 52

MQDTATCSICQGTGFVKTEDNKVRLCECRFKRDVNRELNIPKRYWNANL	100
DTYHPKNVSQNALLTIRVFVHNPNPEEGKGLTFVGSPGVKTHLAVATL	
KAIYEKKGIRGYFFDTKDLIFRLKHMDEGKDTKFLKTVLNSPVLVLDL	200
GSERLSDWQRELISYIITYRYNNLKSTIITTNYSLQREEESSVRISADLA	
SRLGENVVSKIYEMNELLVIKGSDLRKSKKLSTPS	

FIG. 53

ATGAAAAAAGATTGAAAATTGAAGTGGAAAATGTCTCGTTAAAGCCT	100
GGAAATAGATCCCGATGCAGGTGTGGTCTCGTTCCGTGGAAAATTCT	
CCGAAGAGATAGAACACCTTGTGCCTTACTGGAGAAGAACCGCGTTT	200
CGAGTCATCGTAACGGTGTCAAAAAGTAACGGGATCTAAGGGAAA	
GATACTTCCCTCTCAACGGTAATGTGCCTACATAAAAGATGTTGTT	
TCGAAGGAAACAGGCTGATTCTGAAAGTGCTGGAGATTCGCGCGGGAC	300
AGGATCGCCTCCAAACTCAGAACGCAAAAAACAGCTCGATGAACTGCT	
GCCTCCCGAACAGAGATCATGCTGGAGGTTGTGGAGCCTCCCGAACGATC	400
TTTGAAAAAGGAAGTACCAACAACCAGAAAAGAGAGAACGAAACCAAAGGGT	
GAAGAATTGAAGATCGAGGATGAAAACCACATCTTGGACAGAAAACCCAG	500
AAAGATCGTCTTCACCCCCCTCAAAATCTTGAGTACAACAAAAAGACAT	
CGGTGAAGGGCAAGATCTCAAAATAGAGAACGATCGAGGGAAAAGAACG	600
GTCCTTCTGATTTACCTGACAGACGGAGAACGATTCTGATCTGCAAAGT	
CTTCAACGACGTTGAAAAGGTCGAAGGGAAAGTATCGGTGGAGACGTGA	700
TCGTTGCCACAGGAGACCTCCTCTCGAAAACGGGGAGCCCACCCCTTAC	
GTGAAGGGAAATCACAAAACCTCCCGAACGAAAAGGATGGACAAATCTCC	800
GGTTAAGAGGGTGGAGCTCCACGCCATACCAAGTTAGCGATCAGGACG	
CAATAACAGATGTGAACGAATATGTGAAACGAGCCAAGGAATGGGGCTTT	900
CCCGCGATAGCCCTCACGGATCATGGAACGTTAGGCCATACCTTACTT	
CTACGACGCCGAAAGAACGCTGGAATAAGCCCATTTCGGTATCGAAG	1000
CGTATCTGGTGGAGTGACGTGGAGCCCGTCATAAGGAATCTCTCCGACGAT	
TCGACGTTGGAGATGCCACGTTCTCGTCCTCGACTTCGAGACGACGGG	1100
TCTCGACCCGCAGGTGGATGAGATCATCGAGATAGGAGCGGTGAAGATAC	
AGGGTGGCCAGATAGTGGACGAGTACCAACTCTCATAAAGCCTCCAGG	1200
GAGATCTCAAGAAAAAGTTCGGAGATCACCGGAATCACTCAAGAGATGCT	
GGAAAACAAGAGAACGATCGAGGAAGTCTGCCGGAGTTCCTCGGTTTC	1300
TGGAAGATTCCATCATCGTAGCACACAACGCCAACTCGACTACAGATT	
CTGAGGCTGTGGATCAAAAAGTGTGGATTGGACTGGAAAGACCCCTA	1400
CATAGATACGCTGCCCTCGCAAAGTCCCTCTCAAACCTGAGAACGCTACT	
CTCTGGATTCCGTGTGGAAAAGCTCGGATGGGTCCCTCCGGCACAC	1500
AGGGCCCTGGATGACCGAGGGTCACCGCTCAGGTTTCCTCAGGTTCGT	
TGAGATGATGAAGAAGATCGGTATCACGAAGCTTCAGAAATGGAGAAGT	1600
TGAAGGATACGATAGACTACACCGCGTTGAAACCCCTCCACTGCACGATC	
CTCGTTAGAACAAAAAGGGATTGAAAACCTATACAAACTGGTTCTGA	1700
TTCCCTATATAAAAGTACTTCTACGGTGTCCGAGGATCCTCAAAAGTGAGC	
TCATCGAGAACAGAGAACGGACTGCTCGTGGTAGCGCGTGTATCTCCGT	1800
GAGCTCGACGTGCCGCCCTCGAAGGAGCGAGTGTGGACTCGAACCTCGAAGA	
GATCGCGAAGTTCTACGACTACATAGAACGTTAGCTCGACGTTATAG	1900
CCGAAGATGAAGAACGACTAGACAGAGAAAGACTGAAAGAAGTGTACCGA	
AAACTCTACAGAACAGGAAATTGAAACAAAGTTGAGAACGCTCGTCATGACCGG	2000
TGATGTTCATTCCTCGATCCCGAAGATGCCAGGGCAGAGCTGCACTTC	
TGGCACCTCAGGGAAACAGAAAACCTCGAGAACGAGTGTACCGTACCTC	2100
AGAACGACCGAACGAAAGTGTGAGAACGAGTGTACCGTACCTCGAGAGATGA	
AGAGATCGCGAGGGAAAGTCGTGATAGAGAACGAGTGTACCGTACCGATA	2200
TGATCGAGGAAGTGCAGCCGCTCGAGAAAAACTTCACCCGCCGATCATA	
GAGAACGCCGATGAAATAGTGAGAACCTCACCATGAAGCGGGCGTACGA	2300
GATCTACGGTGATCCGCTCCGAAATCGTCCAGAACGCGTGTGGAAAAGG	

FIG. 54A

AACTGAACGCCATCATAAATCATGGATA	CGCCGTTCTCTATCTCATCGCT	2400
CAGGAGCTCGTTCAGAAATCTATGAGC	GATGGTTACGTGGTTGGATCCAG	
AGGATCCGTCGGGCTTCACTCGTGGCA	ATCTCCTCGGAATAACAGAGG	2500
TGAATCCCCTACCACCACATTACAGG	TGTCCAGAGTGC	AAATAACTTGAA
GTGTCGAAGACGACAGATA	CGGAGCGGGTACGAC	2600
CTGTCCAAGATGTGGGGCTCCTCTC	AGAAAAGACGGCACGG	
TTGAAACGTTCATGGGGTCAGGGAG	GGTACAAGGTCCCCGACATAGA	2700
AACTTCTCAGGAGAGTATCAGGAAC	GCTCATCGTTTGTGGAAGAACT	
CTTCGGTAAAGAC	CGACGTCTATAGGGCGGAA	2800
AAAGAAGTGC	GGTACGTGAGAACGCTACGAA	
AAGCTCAGAAAGCGGAAATG	AGAACACTCGTTCCATGATCACGGAGT	2900
GAAGAGAACGACGGTCAGCACCC	AGGGGGCTCATGATCATACCGAAAG	
ACAAAGAAGTCTACGATTCA	CTCCCATA	3000
ACAGCAGGTGTGTT	TCACACGC	
CCTGGTGAAGATA	ACTCGC	
TGCTCAAGGACCTCACCGGAA	ATCGATCCC	
CCCGATACGCTGCCATATT	ATCGAGT	
CGTTGAGCTG	GGAAAGCGATGTGGGAA	3200
CCGAGTTTG	ACGTACGGAA	
GAGCTTGTGAGAATCTCAGGACT	GTGAAACGAGACCAAGAG	3300
CAACGCACGTGATTGGATA	AGTTCGCC	
TCTCGTGTAGGGACGACAT	CGTACGGCTACGG	3400
CCGTCACTGCCTCAAGATCAT	GGCAAGCTCTCG	
CACAGAAGAGATGGAGAGCGAG	AGGTTACGGAA	3500
TCATCGAATCCTG	AGTCAAAATATCTCTCCG	
GTGGCTTACGTGAGTATGGC	AAAGGTACGTT	3600
TCCTCTTCAGTTTACGCGCGT	ACTTCACGATAAAAGGT	
ATCCGGTTCTCGTACTCAGGGAA	AGGTGATCAGTCG	3700
GAAC	ATGCCAAAGACGCCAGAAG	
TGTTCTGGAGGTTGCC	AAAAACGAAAGTGA	3800
CGCCCGACATCTCAAATCCG	ACTGAGAGGTTTCCTC	
TCGCTGAGAATTCCGTT	CTGAGAAATTCTGATAGAAGG	3900
CGAGTCGATAATCAGAGCC	AGGAAAGAAAGCC	
ATCTCATGAAGAGGACCAAGG	TTCAAGGTTCACTTC	4000
AGCCTGGGTGTTCTCGGG	GGGTGGAAG	
C	ACCTCCAGAGACGGAA	4100

FIG. 54B

MKKIENLKWKNVSFKSLEIDPDAGVVLVSVEKFSEEIEDLVRILLEKKTRF	
RVIVNGVQKSNGDLRGKILSLLNGNVPYIKDVFEGNRLLIKVLGDFARD	100
RIASKLRSTKKQLDELLPPGTEIMLEVVEPPEDLLKKEVPQPEKREEPKG	
EELKIEDENHIFGQKPRKIVFTPSKIFEYNKKTsvKGKIFKIEKIEGRT	200
VLLIYLTDGEDSLICKVFNDVEKVEGKSVGDVIVATGDLLLENGETLY	
VKGITKLPEAKRMDKSPVKRVELHAHTKFSDQDAITDVNEYVKRAKEWGF	300
PAIALTDHGKVQAI PYFYDAAKEAGIKPIFGIEAYLVSVDVEPVIRNLSDD	
STFGDATFVVLDFETTGLDPQVDEIIIEIGAVKIQGGQIVDEYHTLIKPSR	400
EISRKSSSEITGITQEMLENKRSIEEVLPFGLFLEDSIIVAHNANFDYRF	
LRLWIKKVMGLDWERPYIDLALAKSLLKLRSYSLDSVVEKGLGPFRHH	500
RALDDARVTAQVFLRFVEMMKKIGITKLSEMEKLDTIDYTALKPFHCTI	
LVQNKKGLKNLYKLVSDSYIKYFYGVPRILKSELIEIREGLLVGSACISG	600
ELGRAALEGASDSELEEIAKFYDYIEVMP LDVIAEDEEDLDRERLKEVYR	
KLYRIAKLNUFVUMTGDVHFLDPEDARGAALLAPQGNRNFENQPALYL	700
RTTEEMLEKAIEIFEDEEIAREVVIENPNRIADMIEEVQPLEKKLHPPII	
ENADEVVRNLTMKRAYEIYGDPLPEIVQKRVEKELNAIINHGYAVLYLIA	800
QELVQKMSDGYVVGSRGSGVSSLVANLLGITEVNPLPPHYRCPECKYFE	
VVEDDRYGAGYDLPNKNCPRCGAPLRKDGHGIPFETFMGFEGDKVPDIDL	900
NFSGEYQERAHRFVEELFGKDHVYRAGTINTIAERSAVGYVRSYEETGK	
KLRKAEMERLVSMITGVKRTTGQHPGLMIIPKDKEVYDFTPIQYPANDR	1000
NAGVFTTHFAYETIHDDLVKIDALGHDDPTFIKMLKDLTGIDPMTIPMDD	
PDTLAIFFSVKPLGVDPVELESDVGTGYGIPFGTEFVRGMLVETRPKSFA	1100
ELVRISGLSHGTDVWLNNARDWINLGYAKLSEVISCRDDIMNFLIHKGME	
PSLAFKIMENVRKKGKITEEMESEMRRLKVPWFIESCKRIKYLFPKAHA	1200
VAYVSMAFRIAYFKVHYPLQFYAAYFTIKGDQFDPVLVLRGKEAIKRRLR	
ELKAMPACKDAQKKNEVSVLEVALEMILRGFSFLPPDIFKSDAKKFLIEGN	1300
SLRIPFNKLPGLGDSVAESIIRAREEKPFTSVEDLMKRTKVNKNHIELMK	
SLGVLGDLPETEQFTLF	1367

FIG. 55

GTGCTGCCATGATATGGAACGACACC GTTT GCGTCGTAGACACAGA
AACCA CGGGAA CCGATCCCTTGC CGGAGACCGGATAGTTGAAATAGCCG 100
CTGTTCTGTCTCAAGGGGAAGATCTACAGAAACAAAGCGTTCACTCT
CTCGTGAATCCCAGAATAAGAATCCCTGC CGT GATT CAGAAAGTTCACCG 200
TATCAGCAACATGGACATCGT GGAAGCGCCAGACATGGACACAGTTACG
ATCTTT CAGGGATTACGTGAAGGGAACGGTGCTCGT GTT CACAACGCC 300
AACTTCGACCTCACTTTCTGGATATGATGGCAAAGGAAACGGGAAACTT
TCCAATAACGAATCCCTACATCGACACACTCGATCTTCAGAAGAGATCT 400
TTGGAAGGCCTCATTCTCTCAAATGGCTCTCCGAAAGACTT GGAATAAAA
ACCACGATA CGGCACCGT GCTCTCCAGATGCCCTGGTACCGCAAGAGT 500
TTTTGTGAAGCTT GTT GAATTCTTGGT GAAAACAGGGTCAACGAATTCA
TACGTGGAAAACGGGGG 567

FIG. 56

MLAMIWNDTVFCVVDTE TTGTD PFAGDRIVEIAAVPVFKGKIYRNKA FHS
LVNPRIPIALIQKVHG ISNMD IVEAPDMDTVYDLFRDVKGTVL VFHNA 100
NFDLTFIDMMAKETGNFPITNPYIDLDLSEEIFGRPHSLKWL SERLGIK
TTIRHRALPDALVTARVFVKLVEFLGENRVNEFIRGKRG 189

FIG. 57

GTGGAAGTTCTTACAGGAAGTACAGGCCAAGACTTTCTGAGGGTTGT	
CAATCAGGATCATGTGAAGAAGGCATAATCGGTGCTATTAGAACAGAAACA	100
GCGTGGCCCACGGATACATATTGCCGGTCCGAGGGAACGGGAAGACT	
ACTCTTGCCAGAATTCTCGCAAATCCCTGAAGTGTGAGAACAGAAAGGG	200
AGTTGAACCTGCAATTCTGCAGAGCCTGCAGAGAGATAGACGAGGGAA	
CCTTCATGGACGTGATAGAGCTGACGCCCTCAACAGAGGAATAGAC	300
GAGATCAGAAGAATCAGAGACGCCGTTGGATACAGGCCATGGAAGGTAA	
ATACAAAGTCTACATAATAGACGAAGTTACATGCTCACGAAAGAACGCT	400
TCAACCGCCTCCTCAAAACACTCGAACGAAACCTCCTCCACGTCGTGTT	
GTGCTGGCAACGACAAACCTTGAGAAGGTTCTCCACGATTATCTCGAG	500
ATGTCAGGTTTCGAGTTCAAGAACATTCCGACGGCTCATCGAAAAGA	
GGCTCCAGGAAGTTGCGGAGGCTGAAGGAATAGAGATAGACAGGGAAAGCT	600
CTGAGCTTCATCGCAAAAGAGCCTCTGGAGGCTTGAGAGACGCGCTCAC	
CATGCTCGAGCAGGTGTTGAGAAGGAAAGATAGATCTCGAGA	700
CGGTACACAGGGCCTGGGTTGATACCGATAACAGGTTGTCGCGATTAC	
GTGAAACGCTATCTTCTGGTGTGAAAAGGGTCTCACCGTTCTCGA	800
CGACGTTCTATTACAGCGGAAGGACTACGAGGTGCTCATTAGGAAGCAG	
TCGAGGATCTGGTCGAAGACCTGGAAAGGGAGAGAGGGGTTTACAGGTT	900
TCAGCGAACGATATAGTCAGGTTTCGAGACAACCTCTGAATCTCTGAG	
AGAGATAAAAGTCGCCGAAGAAAAACGACTCGTCTGAAAGTGGGTTCGG	1000
CTTACATAGCGACGAGGTTCTCACCACAAACGTCAGGAAAACGATGTC	
AGAGAAAAAAACGATAATTCAAATGTACAGCAGAAAGAAGAGAAGAAAAGA	1100
AACGGTGAAGGCAAAAGAAGAAAAACAGGAAGACAGCGAGTTGAGAAC	
GCTTCAAAGAACTCATGGAAGAACTGAAAGAAAAGGGCAGTCCTCTATC	1200
TTTGTGCTCTCAGCCTCTCAGAGGTGCAAGTTGACGGAGAAAAGGTGAT	
TATTCTTTGATTCATGAAAGCTATGCATTACGAGTTGATGAAGAAAA	1300
AACTGCTGAGCTGGAAAACATTCTAGAAAACCTGGAAAAAGTA	
GAAGTTGAACCTCGACTGATGGGAAAAGAAGAAACAATCGAGAAGGTTTC	1400
TCAGAAGATCCTGAGATTGTTGAACAGGGAGGGAA	

FIG. 58

MEVLYRKYPKTFSEVVNQDHVKKAIIGAIQKNSVAHGYIFAGPRGTGKT	
TLARILAKSLNCENRKGVEPCNSCRACREIDEFTFMDVIELDAASNRGID	100
EIRRIRDAGYRPMEGKYKVIIDEVHMLTKEAFNALLKLEEPSSHVVF	
VLATTNLEKVPPTIISRCQVFERNIPDELIKRLQEVAEAEGLIEDREA	200
LSFIAKRASGGLRDALTMLEQVWKFSEGKIDLETVHRALGLIPIQVVRDY	
VNAIFSGDVKRVFTVLDDVYYSGKDYEVLIQEAVEDLVEDLERERGVYQV	300
SANDIVQVSQRQLNLLREIKFAEKRLVCKVGSAYIATRFSTTNVQENDV	
REKNDNSNVQQKEEKETVKAKEEKQEDSEFEKRFKELMEELKEKGDLISI	400
FVALSLSEVQFDGEKVIISFDSSKAMHYELMKKLPELENIFSRKLGKKV	
EVELRLMGKEETIEKVSKILRLFEQEG	478

FIG. 59

ATGAAAGTAACCGTCACGACTCTTGAATTGAAAGACAAAATAACCATCGC	100
CTCAAAAGCGCTCGCAAAGAAATCCGTGAAACCCATTCTGCTGGATTTC	
TTTTCGAAGTGAAGAGATGAAATTCTACATCTGCACGATCTCGAG	200
ACCGGAGTCAAAGCAACCGTGAATGCCGCTGAAATCTCCGGTGAGGCACG	
TTTTGTGGTACCAAGGAGATGTCATTAGAAGATGGTCAAGGTTCTCCAG	
ATGAGATAACGGAACCTTCTTAGAGGGGATGCTCTGTTATAAGTCT	300
GGAAGCACCCTTTCAGGATCACCACCATGCCCGCGACGAATTCCAGA	
GATAACGCCTGCCGAGTCCTGGAATAACCTCGAAGTTGACACTTCGCTCC	400
TCGAGGAATGGTTGAAAGGTATCTCGCCGCTGCCAAAGACGAGTTC	
ATGCGAAATCTGAATGGAGTTCTGGAAACTCCACAAGAATCTCTCAG	500
GCTGGTTGCAAGTGTGGTTTCAGACTTGCACCTGCTGAAGAGCAGATAG	
AAAACGAGGAAGAGGGAGTTCTGCTCTTGAAGAGCATGAAAGAA	600
GTTCAAAACGTGCTGGACAACACAAACGGAGCCGACTATAACGGTGAGGTA	
CGATGGAAGAAGGGTTCTCTGTGACAAATGATGTAGAAACGGTGATGA	700
GAGTGGTCGACGCTGAATTCCGATTACAAAAGGGTATCCCCGAAACT	
TTCAAAACGAAAGGGTTCCAGAAAAGAACTCAGGAAATCTTGAA	800
GAGGGTATGGTGATTGCCAGCAAGGGAAAGCGAGTCGTGAAGTTGAAA	
TAGAAGAAAACGTTATGAGACTTGTGAGCAAGAGCCGGATTATGGAGAA	900
GTGGTCGATGAAGTTCAAAAAGAAGGGAAAGATCTGTGATCGC	
TTTCAACCGAAGTTCATCGAGGACGTTGAAGCACATTGAGACTGAAG	1000
AAATCGAAATGAACCTCGTTGATTCTACCACTGTCAGATAAATCCA	
CTCGATATTCTGGATACCTTACATAGTGTGATGCCATCAGACTGGCA	1098

FIG. 60

MKVTVTLELKDKITIASAKALAKSVKPILAGFLFEVKDGNFYICATDLE	100
TGVKATVNAAEISGEARFVVPGDVIQKMKVVLPEDEITELSLEGDALVISS	
GSTVFRITTMPADEFPEITPAESGITFEVDTSLLLEEMVEKVIIFAAAKDEF	200
MRNLNGVFWELHKNLLRLVASDGFRLLALAEEQIENEESASFLSLKSMKE	
VQNVLNDNTTEPTITVRYDGRRLSLSNDVETVMRVVDAEFPDYKRVIPET	300
FKTKVVVSRKELRESLKRVMVIASKGSESVKFEIEENVMRLVSKSPDYGE	
VVDEVEVQKEGEDLVIAFNPKFIEDVLKHIEETEEIEMNFVDSTSPCQINP	366
LDISGYLYIVMPIRLA	

FIG. 61

ATGCCAGTCACGTTCTCACAGGTACTGCAGAAACTCAGAAGGAAGAATT	
GATAAAGAAACTCCTGAAGGATGGTAACGTGGAGTACATAAGGATCCATC	100
CGGAGGATCCGACAAGATCGATTCATAAGGTCTTACTCAGGACAAG	
ACGATCTTCAACAAGACGATCATGACATCGTCAATTGATGAGTG	200
GAAAGCACAGGAGCAGAACGCTCTCGTTGAACCTTGAAAACGTACCGG	
AAGACGTTCATATCTCATCCGTTCTCAAAAAACAGGTGGAAAGGGAGTA	300
GCGCTGGAGCTTCCGAAGCCATGGGAAACGGACAAGTGGCTTGAGTGGAT	
AGAAAAGCGCTTCAGGGAGAATGGTTGCTCATCGATAAAGATGCCCTC	400
AGCTGTTTTCTCCAAGGTTGGAACGAACGACCTGATCATAGAAAGGGAG	
ATTGAAAAACTGAAAGCTTATTCCGAGGACAGAAAGATAACGGTAGAAGA	500
CGTGGAAAGAGGTGTTTACCTATCAGACTCCGGATACGATGATTTT	
GCTTGCTGTTCCGAAGGAAAAGGAAGCTCGCTCACTCTCTGTGCG	600
CAGCTGTGAAAACCACAGACTCCGTGGTGAATTGCCACTGTCTTGC	
TCACTTCTGGATCTCTCAAATCCTCGTTCTGTGACAAAGAAAAGAT	700
ACTACACCTGGCCTGATGTGTCCAGGGTGTCAAAGAGCTGGAAATTCCC	
GTTCCCTCGTGTGGCTCGTTCTCGGTTCTCCTTTAAGACCTGGAAATT	800
CAAGGTGATGAACCACCTCCTACTACGATGTGAAGAAGGTTAGAAAGA	
TACTGAGGGATCTCTACGATCTGGACAGAGCCGTGAAAAGCGAAGAAGAT	900
CACAAACCGTTCTCCACGAGTTCATAGAAGAGGTGGCACTGGATGTATA	
TTCTCTTCAGAGAGATGAAGAA	972

FIG. 62

MPVTFLTGTAEQKEELIKLLDGNVEYIRIHPEDPDKIDFIRSLLRTK	
TIFSNKTIIDIVNFDEWKAQEOKRIVELLKNVPEDVHIFIRSQKTGGKGV	100
ALELPKPWETDKWLEWIEKRFRENGLLIDKDALQLFSKVGTNDLIIERE	
IEKLKAYSEDRKITVEDVEEVFTYQTPGYDDFCFAVSEGKRKLAHSLLS	200
QLWKTTESVVIATVLANHFSDLFKILVLVTKKRYTWPDVSRVSKELGIP	
VPRVARFLGSFKTWKFVNMHLLYYDVKKVRKILRDLYDLDRAVKSEED	300
PKPFFHEFIEEVALDVYSLQRDEE	

FIG. 63

ATGAAACGATTTGATCAGAAAGTACGCTAAAGATCAACTGGAAACTTGAA	
AAGGATCATAGAAAAGTCTGAAGGAATATCCATCCTCATAAATGGAGAAG	100
ATCTCTCGTATCCGAGAGAAGTATCCCTGAACTTCCCGAGTACGTGGAG	
AAATTCCCCGAAGGCCTCGGATGTTCTGGAGATAGATCCCGAGGGGGA	200
GAACATAGGCATAGACGACATCAGAACGATAAAGGACTTCCTGAAC	
CCCCGAGCTACACGAGAAAGTACGTGATAGTCCACGACTGTGAAAGA	300
ATGACCCAGCAGGCCGAACCGCTTCTGAAGGCCCTGAAGAACACC	
AGAATACGCTGTGATCGTTCTGAACACTCGCCGCTGGCATTATCTACTGC	400
CGACGATAAAAGAGCCGAGTGTTCAGAGTGGTTGTGAACTTCCAAAGGAG	
TTCAGAGATCTCGTCAAAGAGAAAATAGGAGATCTCTGGGAGGAAC	500
ACTTCTTGAGAGAGACTTCAAAACGGCTCTGAAGCCTACAAACTTGGTG	
CGGAAAAACTTCTGGATTGATGAAAGTCTCAAAGTTGGAGACGGAA	600
AAACTCTGAAAAAGGTCTTCAAAAGGCCTCGAAGGTTATCTCGCATG	
TAGGGAGCTCCTGGAGAGATTTCAAAGGTGGAATCGAAGGAATTCTTG	700
CGCTTTTGATCAGGTGACTAACACGATAACAGGAAAAGACCGCTTCTT	
TTGATCCAGAGACTGACAAGAATCATTCTCACGAAAACACATGGGAAAG	
CGTTGAAGATCAAAAAGCGTGTCTTCCTCGATTCAATTCTCAGGGTGA	
AGATAGCGAATCTGAACACAACACTCACTCTGATGAAACATCCTCGCGATA	
CACAGAGAGAGAAAGAGAGGGTGTCAACGCTGGAGC	900

FIG. 64

MNDLIRKYAKDQLETLKRIIEKSEGISILINGEDLSYPREVSLELP	EYVE
KFPPKASDVLEIDPEGENIGIDDIRTIKDFLNYSPELYTRKYVIVHDCER	100
MTQQAANANFLKALEEPPEYAVIVLNRRWHYLLPTIKSRVFRVVNPKE	
FRDLVKEKIGDLWEELPLLERDFKTALEAYKLGAEKLSGLMESLK	200
VLETE	
KLLKKVLSKGLEGYLACRELLERFSKVESKEFFALFDQVTNTITGKD	
AF	
LIQRRLTRIILHENTWESVEDKSVSFLDSILRVKIANLNNKL	300
LMNIALIH	
RERKRGVNAWS	

FIG. 65

ATGTCTTCTCAACAAGATCATACTCATAGGAAGACTCGTGAGAGATCC
CGAAGAGAGATAACCGCTCAGCGGAACCTCAGTCACCACCTCACCATAG 100
CGGTGGACAGGGTTCCCAGAAAGAACGCGCCGGACGACGCTCAAACGACT
GATTTCCTCAGGATCGTCACCTTCCAAGACTGGCAGAGTCGCTAGAAC 200
CTATCTCACCAAGGAAGGCTCGTCTCGTGAAGGTGAAATGAGAATGA
GAAGATGGAAACACCCACTGGAGAAAAGAGGGTATCTCCGGAGGTTGTC 300
GCAAACGTTGTTAGATTATGGACAGAAAAACCTGCTGAAACAGTTAGCGA
GACTGAAGAGGAGCTGGAAATACCGGAAGAAGACTTTCCAGCATACT 400
TCAGTGAAGATGAACCACCATTT

FIG. 66

MSFFNKKIILIGRLVRDPEERYTLSGTPVTTFTIAVDRVPRKNAPDDAQTT
DFFRIVTFGRLAEFARTYLTGRLVLVEGEMRMRRWETPTGEKRVSPVV 100
ANVVRFMDRKPAETVSEEELEIPEEDFSSDTFSEDEPPF

FIG. 67

ATGCGTGTCCCCGCACAACCTAGAGGCCGAAGTTGCTGTGCTCGGAAG	100
CATATTGATAGATCCGTCGGTAATAAACGACGTTCTGAAATTGAGCC	
ACGAAGATTCTATCTGAAAAAACACCAACACATCTCAGAGCGATGGAA	200
GAGCTTACGACGAAGGAAAACCGGTGGACGTGGTTCCGTCGTGACAA	
GCTTCAAAGCATGGAAAATCGAGGAAGTAGGTGGAGATCTGAAAGTGG	
CCCAGCTCGCTGAGGCTGTGCCAGTTCTGCACACGCACTCACTACCG	300
GAGATCGTCAAGGAAAATCCATTCTGAGGAAGACTCATTGAGATCTCCAG	
AAAAATCTCAGAAAGTGCCTACATGGAAGAAGATGTGGAGATCCTGCTCG	400
ACAAACGAGAAAAGATGATCTCGAGATCTCAGAGATGAAAACGACAAA	
TCCTACGATCATCTGAGAGGCATCATGCACCGGGTGGAAACCTGGAA	500
GAACCTCAGGGAAAGAGCCAACCTTATAGAACCCGGTGTGCTCATACGG	
GAECTACCAACGGGATTCAAAAGTCTGGACAAACAGACCACAGGGTCCAC	600
AGCTCCGATCTGGTATAATAGCAGCAGACCCCTCCATGGGAAAACCTC	
CTTCGCACTCTCAATAGCAGGAACATGGCTGTCAATTTCGAAATCCCCG	700
TCGGAATATTCACTCGAGATGTCCAAGGAACAGCTCGCTCAAAGACTA	
CTCAGCATGGAGTCCGGTGTGGATCTTACAGCATCAGAACAGGATACT	800
GGATCAGGAGAAGTGGAAAGACTCACAAATAGCGGTTCTAAACTCTACA	
AAGCACCCATAGTTGTGGACGATGAGTCACTCCTCGATCCGCGATCGTTG	900
AGGGCAAAAGCGAGAAGGATGAAAAAGAACATACGATGTAAAAGCCATT	
TGTCGACTATCTCCAGCTCATGCACCTGAAAGGAAGAAAAGAACAGAC	1000
AGCAGGAGATATCCGAGATCTCGAGATCTCTGAAGCTCCTGCGAGGGAA	
CTCGACATAGTGGTGTAGCGCTTCACAGCTTCGAGGGCCGTAGAACAA	1100
GAGAGAACAAAAGACCGAGGCTGAGTGACCTCAGGGAATCCGGTGC	
AGAACAGGACGCAGACACAGTCATCTCATCTACAGGGAGGAATTAC	1200
AGGAGCAAAATCCAAAGAGGAAAGCAAGCTTACGAACCTCACGAAGC	
TGAAATCATATAAGTAAACAGAGAAACGGTCCGTTGGAACGATCACTC	1300
TGATCTCGACCCAGAACGGTTACGTTCCATGAAGTCGATGTGGTGCAT	
TCA	1353

FIG. 68

MRVPPHNLEAEAVLGSILIDPSVINDVLEILSHEDFYLKKHQHIFRAME	100
ELYDEGPVDVSVCDKLQSMGKLEEVGGDLEVAQLAEAVPSSAHALHYA	
EIVKEKSILRKLIIEISRKISESAYMEEDVEIILDNAEKMIFEISEMKTK	200
SYDHLRGIMHRVFENLENFRERANLIEPGVLITGLPTGFKSLDKQTTGFH	
SSDLVIIAARPSMGKTSFALSIARNMAVNFEIPVGIFSLEMSKEQLAQRL	300
LSMESGVDLYSIRTGYLDQEKGWERLTIAASKLYKAPIVVDDESLLDPRSL	
RAKARRMKKEYDVKAIFVDYLQLMHLKGRKESRQQEISEISRSLKLLARE	400
LDIVVIALSQLSRAVEQREDKRPRLSLDLRESGAIEQDADTVIFIYREYY	
RSKKSKEEKLHEPHEAEIIIGKQRNGPVGTITLIFDPRTVTFHEVDVHS	451

FIG. 69

GTGATTCTCGAGAGGTATCGAGGAAATAAAGAAAAGGTTGACATCGT	1 00
AGAGGTCACTCCGAGTACGTGAATCTTACCCGGTAGGTTCCCTACA	
GGGCTCTCTGCCCCCTTCATTAGAAACCAATCCTCTTCTACGTTCAT	2 00
CCGGGTTGAAGATATACCATTGTTGGCTCGCGTGCAGTGGAGACGT	
CATCAAATTCAGAAATGGAAGGGATCAGTTCCAGGAAGCGCTGG	3 00
AAAGACTTGCCAAAAGAGCTGGGATTGATCTTCTCTACAGAACAGAA	
GGGACTTCTGAATACGGAAAATACATTGTTGTACCGAAGAAACGTGGAA	4 00
AAGGTACGTCAAAGAGCTGGAGAAATCGAAAGAGGAAAAGACTATTTAA	
AAAGCAGAGGCTCTCTGAAGAAGATATAGCAAAGTCGGCTTGGGTAC	5 00
GTCCCCAAGAGATCCAGCATCTCTATAGAAGTGCAGAAGGCATGAACAT	
AAACACTGGAAGAACCTTGTCAAGATACGGTATCGCGCTAAAAAGGGTGATC	6 00
GATTGTTGATAGATTGAAGGAAGAATCGTTCCAATAAAAGAACGAC	
AGTGGTCATATTGGCTTTGGTGGCGTCTCGGCAACGAAGAAC	7 00
GAAGTATTGAACTCTCAGAGACCAAGGTATTTTCAAGAAGAACGACCC	
TTTTCTCTTCGATGAGGCGAAAAAAGTGGCAAAGAGGTTGGTTTTTC	8 00
GTCATCACCGAAGGCTACTTCGACCGCCTCGCATTCAAGAACGATGGAAT	
ACCAACGGCGGTGCTGTTCTGGGCGAGTCTTCAAGAGAGGCGATTC	9 00
TAAAACCTTCGGCGTATTGAAACGTCAACTGTGTTCGATAATGAC	
AAAGCAGGCTTCAGAGCCACTCTCAAATCCCTGAGGATCTCTAGACTA	1 000
CGAATTCAACGTGTTGTGCAACCCCTCTCCTACAAAGACCCAGATG	
AACTCTTCAGAAAGAAGGAGAAGGTTCATTGAAAAAGATGCTGAAAAC	1 100
TCGCGTTGTTGAAATATTCTGGTACGGCTGGTGGAGGTCTTCTTGA	
CAGGAACAGCCCCGGGTGTGAGATCCTACCTTCTTCTCAAAGGTT	1 200
GGGTCCAAAAGATGAGAAGGAAGGATATTGAAACACATAGAAAATCTC	
GTGAATGAGGTTTACGATGAGGGAGAGGACTGGCTTATTGTTTGAAC	1 300
CTTTTTGAAAGCGACAGGTCTAACACTATGCCTGTTCATGAGACCAAGT	
CGTCAAAGGTTACGATGAGGGAGAGGACTGGCTTATTGTTTGAAC	1 400
TACGAGGATTGAGGGAAAAGATTCTGGAACTGGACTTAGAGGTTACTGGA	
AGATAAAAACGCAGGGAGTTTCAAGAGAGTCTCACTGGAGAACGATT	1 500
TGAACAAAGTCATAGAAAACCTCCAAAAGAGCTGAAAGACTGGATT	
GAGACAATAGAAAGCATTCCCTCAAAGGATCCGAGAAATTCCCTCGG	1 600
TGACCTCTCGAAAAGTTGAAAATCCGACGGATAGAGAGACGTATCGCAG	
AAATAGATGATATGATAAAGAAAGCTTCAAACGATGAAGAACGGTCTT	1 695
CTTCTCTATGAAAGTGGATCTCCTCAGAAAATAAGAGGAGG	

FIG. 70

MIPREVIEEIKEKVDIVEVISEYVNLTRVGSSYRALCPFHSETNPSFYVH	
PGLKIYHCFGCGASGDVIKFLQEMEGISFQEALERLAKRAGIDLSLYRTE	100
GTSEYGYKIRLYEETWKRYVKELEKSKEAKDYLKSRGFSEEDIAKFGFGY	
VPKRSSISIEVAEGMNTILEELVRYGIALKKGDRFVDRFEGRIVVPIKND	200
SGHIVAFGGRALGNEEPKYLNSPETRYFSKKTLFLFDEAKKVAKEVGFF	
VITEGYFDALAFRKDGIPAVAVILGASLSREAILKLSAYSKNVILCFND	300
KAGFRATLKSLEDLLDYEFNVLVATPSPYKDPDELFKGEGEGLKKMLKN	
SRSFEYFLVTAGEVFFDRNSPAGVRSYLSFLKGWVQKMRRKGYLKHIENL	400
VNEVSSLQLIPENQILNFFESDRSNTMPVHETKSSKVYDEGRGLAYLFLN	
YEDLREKILELDLEVLEDKNAREFFKRVSLGEDLNKVIENFPKELKDWF	500
ETIESIPPPKDPEKFLGDLSEKLKIRRIERIAEIDDMIKKASNDEERRL	
LLSMKV DLLRKIKRR	565

FIG. 71

ATGGCTCTACACCCGGCTCACCTGGGGCAATAATCGGGCACGAGGCCGT	
TCTCGCCCTCCTCCCCGCCTCACCGCCCAGACCCCTGCTCTTCTCCGGCC	100
CCGAGGGGGTGGGGCGGCCACCGTGGCCCGCTGGTACGCCTGGGGGCTC	
AACCGCGGCTTCCCCCGCCCTCCCTGGGGAGCACCCGGACGTCTCGA	200
GGTGGGGCCCAAGGCCGGGACCTCCGGGGCCGGAGGTGCGGCTGG	
AGGAGGTGGCCATCCTGGACTCGGCCACCTCCTCACCGAGGCCGCGC	300
GTGAAGGTGGCATCGGGACTCGGCCACCTCCTCACCGAGGCCGCGC	
CAACGCCCTCCTCAAGCTCCTGGAGGAGCCCCCTTCTACGCCGCATCG	400
TCCTCATGCCCTAACGCCGCCACCCCTCCCTCCCCACCTGGCCTCCGG	
GCCACGGAGGTGGCATTGCCCTCGCTACGCCGCCGGGCCCCGGGCC	500
CACCCAGGACCCGGAGCTCCTCGCTACGCCGCCGGGCCCCGGGCC	
TCCTTAGGGCCCTCCAGGACCCGGAGGGTACCGGGCCGCATGGCCAGG	600
GCGCAAAGGGTCTGAAAGCCCCGCCCTGGAGCGCCTCGCTTGCTTCG	
GGAGCTTTGGCCGAGGAGGAGGGGTCCACGCCCTCCACGCCGTCTAA	700
AGGCCCGGAGCACCTCCTGGCCCTGGAGCGGGCGCGGGAGGCCCTGGAG	
GGGTACGTGAGCCCCGAGCTGGCCTCGCCGGCTGGCCTAGACTTAGA	800
GACA	

FIG. 72

MALHPAHPGAIIGHEAVLALLPRITAQTLLFSGPEGVGRRTVARWYAWGL	
NRGFPPPSLGEHPDVLEVGPKARDLRGRAEVRLVEVAPLLEWCSSHPRER	100
VKVAILDSAHLTEAAANALLKLLEEPSYARIVLIAPS RATLLPTLASR	
ATEVAFAPVPEEALRALTQDPPELLRYAAGAPGRLLRALQDPEGYRARMAR	200
AQRVLKAPPLERLALLRELLAEEGVHALHAVLKRPEHLLALERAREALE	
GYVSPELVLARLALDLET	268

FIG. 73

ATGCTGGACCTGAGGGAGGTGGGGGAGGC GGAGTGGAAAGGCCCTAAAGCC	100
CCTTTTGGAAAGCGTGCCCGAGGGCGTCCCCGTCCCTCCTCCTGGACCCCTA	
AGCCAAGCCCCTCCCGGGCGGCTTCTACCGGAACCGGGAAAGGCGGGAC	200
TTCCCCACCCCCAAGGGGAAGGACCTGGTGCGGCACCTGGAAAACCGGGC	
CAAGCGCCTGGGCTCAGGCTCCCGGGCGGGTGGCCCAGTACCTGGCCT	
CCCTGGAGGGGACCTCGAGGCCCTGGAGCGGGAGCTGGAGAACGCTGCC	300
CTCCTCTCCCCACCCCTCACCCCTGGAGAACGGTGGAGAACGGTGGTGGCCCT	
GAGGCCCCCCCTCACGGGCTTGAACCTGGTGCGCTCCGTCTGGAGAACGG	400
ACCCCAAGGAGGCCCTCGGCCTAGGC GGCTCAAGGAGGAGGGGAG	
GAGCCCTCAGGCTCCTCGGGCCCTCTCCCTGGCAGTTGCCCTCCTCGC	500
CGGGGCTTCTTCCTCCCTCCGGAAAACCCAGGCCAAGGAGGAGGACC	
TCGCCCCGCTCGAGGCCACCCCTACGCCGCCGCCGCGCCCTGGAGGCG	600
GCGAACGCGCTCACGGAAGGCCCTCAAGGAGGCCCTGGACGCCCTCAT	
GGAGGCGGAAAAGAGGCCAAGGGGGAAAGACCCGTGGCTGCCCTGG	700
AGGCGGGTCCCGCCTCGCCCTTGA	

FIG. 74

MVIAFTGDPFLAREALLEEARLRGLSRFTEPTPEALAQALAPGLFGGGGA	
MLDLREVGEAEWKALKPLLESVPEGVPVLLDPKPSPSRAAFYRNRRD	100
FPTPKGKDLVRHLENRAKRLGLRLPGGVAQYLASLEGDLEALERELEKLA	
LLSPPLTLEKVEKVVALRPPLTGFDLVRSVLEKDPEAKLRLGGLKEEGE	200
EPLRLIGALSWQFALLARAFFLLRENPRPKEEDLARLEAHPYAARRALEA	
AKRLTEEALKEALDALMEAEKRAKGGKDPWLALEAAVLRLAR	292

FIG. 75

ATGGCTCGAGGCCTGAACCGCGTTTCCCTCATCGGCGCCCTGCCACCCG
 100
 GCCGGACATGCCTACACCCCGGGGGCTGCCATTGGACCTGACCC
 TCGCCGGTCAGGACCTGCTTCCATAACGGGGGGAACCGGAGGTG
 200
 TCCTGGTACCAACGGGTGAGGCTCTTAGGCCAGGCGGAGATGTGGG
 CGACCTCTGGACCAAGGGCAGCTCGTCTCGTGGAGGGCCCTGGAGT
 ACCGCCAGTGGAAAGGGAGGGGAGAACGGGAGCGAGCTCAGATCCG
 300
 GCCGACTTCCGGACCCCTGGACGACCGGGGAAGAAGCGGGCGGAGGAC
 AGCCGGGCCAGCCCAGGCTCCGCCGCCCTGAACCAGGTCTCCTCAT
 400
 GGGCAACCTGACCCGGGACCCCGAACCTCCCTACACCCCCCAGGGCACCG
 CGGTGGCCCGCTGGCCTGGCGGTGAACGAGCGCCGCCAGGGGGCGGAG
 500
 GAGCGCACCCACTTCGTGGAGGTTCAAGGCCTGGCGCACTGGCGGAGTG
 GGCGCCGAGCTGAGGAAGGGCAGGGCTTTCTGTGATCGGCAGGTGG
 TGAACGACTCCTGGACAGCTCCAGCGGCAGCGGCCTTCAGACCCGT
 GTGGAGGCCCTCAGGCTGGAGCGCCCAACCGTGGACCTGCCAGGCCTG
 CCCAGGCCGGCGAACAGGTCCCAGAAGTCCAGACGGGTGGGTGGACA
 TTGACGAAGGCTTGGAAAGACTTCCGCCGGAGGAGGATTGCCGTTTGA
 600
 700
 800
 GCACGAA

FIG. 76

MARGLNRVFLIGALATRPDMRYTPAGLAILDLTLAGQDLLSDNGGEPEV
 SWYHRVRLLGRQAEMWGDLLDQGQLVFVEGRLEYRQWEREGEKRSELQIR
 ADFLDPLDRGKRAEDSRGQPRLRAALNQVFLMGNLTRDPELRYTPQGT
 AVARLGLAVNERRQAEERTHFVEVQAWRDLAEWAAELRKGDGLFVIIGRL
 VNDSWTSSSGERRFQTRVEALRLERPTRGPAQACPGRNRREVQTGGVD
 IDEGLEDFPPEEDLPF
 100
 200
 266

FIG. 77

AATTCCGACATTCATTGAATCGTTATTCCGCTTAAAAAGAAGGCAA
 GTTGCTCGTTGATGTGAAAAGACCGGGGAGCATCGTACTGCAGGCGCGCT 100
 TTTTCTCTGAAATCGTAAAAAACTGCCAACAAACGGTGGAAATCGAA
 ACGGAAGACAACCTTTGACGATCATCCGCTCGGGCACTCAGAATTCCG 200
 CCTCAATGGGCTAACGCCGACGAATATCCGCGCTGCCGCAAATTGAAG
 AAGAAAACGTGTTCAAATCCCCTGGTATTGAAAACCGTGATTGG 300
 CAAACGGTGTCCCGTTCTACATCGGAAACCGGCCAATCTTGACAGG
 TGTCAACTGGAAAAGTTGAACATGGCGAGCTTGTCTGCACAGCGACCGACA 400
 GTCATCGCTTAGCCATGCCAAAGTAAAATTGAGTCGAAAATGAAGTA
 TCATACAAACGTCGTATCCCTGGAAAAGTCTTAATGAGCTCAGCAAAAT 500
 TTGGATGACGGCAACCACCCGGTGGACATCGTCATGACAGCCAATCAAG
 TGCTATTTAAGGCCGAGCACCTCTCTCTTCCCCTGGCTGTTGACGGC 600
 AACTATCCGGAGACGGCCGCTTGATTCCAACAGAAAGCAAAACGACCAT 700
 GATCGTCATGCAAAAGAGTTCTGCAGGCAATCGACCGAGCGTCTTGCA
 TTGCTCGAGAAGGAAGAACACGTTGTGAAACTGACGACGCTTCTGGAG
 GGAATGTCGAAATTCTCGATTCTCGAGATCGGGAAAGTGAACATTGTTCA
 CAGCTGAAACGGAGTCTCTGAAGGGGAAGAGTTGAACATTGTTCA
 CGCGAAATATATGATGGACCGCGTGGCGCTTGATGGAACAGACATT
 CAAATCAGCTTCACTGGGCCATGCCGCTTCTGTTGCGCCGCTTC
 ACCGATTGATGCTCAGCTCATTTGCCGGTGAGAACATAT 900
 992

FIG. 78

NSDISIIESFIPLEKEGKLLVDVKRPGSIVLQARFFSEIVKKLPQQTVEI
 ETEDNFLTIIRSGHSEFRILNGLNADEYPRLPQIEEENVFQIPADLLKTVI 100
 RQTVFAVSTSETRPILTGVNWKVEHGEVCTATDSHRLAMRKVKIIIESEN
 EVSYNVVI PGKSLNELSKII LDGNHPVDIVMTANQVLFKAEHLLFFSRL 200
 LDGNYPETARLIPTESKTTMIVNAKEFLQAIIDRASLLAREGRNNVVKLTT
 LPGGMLEISSISPEIGKVTEQLQTESLEGEELNISFSAKYMMDALRALDG 300
 TDIQISFTGAMRPFLRLPLHTDSMLQLILPVRTY

FIG. 79

ATGATTAAACCGCGTCATTTGGTCGGCAGGTTAACGAGAGATCCGGAGTT
GCCTTACACTCCAAGCGGAGTGGCTGTTGCCACGTTACGCTCGCGGTCA 100
ACCGTCCTTACAAATCAGCAGGGCGAGCGGGAAACGGATTTTATTCAA
TGTGTCGTTGGCGCCAGGCGAAAACGTCGCCAACTTTTGAAAAA 200
GGGGAGCTTGGCTGGTGTGATGGCGACTGCAAACCCGCAGCTATGAAA
ATCAAGAAGGTGGCGTGTGTACGTGACGGAAGTGGTGGCTGATAGCGTC 300
CAATTCTTGAGCCGAAAGGAACGAGCGAGCAGCGAGGGCGACAGCAGG
CGGCTACTATGGGGATCCATTCCCATTGGGCAAGATCAGAACCAAT 400
ATCCGAACGAAAAAGGGTTGGCCCATCGATGACGATCCTTCGCCAAT
GACGGCCAGCCATCGATATTCTGATGATGATTGCCGTTT 492

FIG. 80

MINRVILVGRLTRDPELRYTPSGVAVATFTLAVNRPFTNQSYENQEGRV
YVTEVVADSVQFLEPKGTSEQRGATAGGYYQGERETDFIQCVVWRRQAEN 100
VANFLKKGSLAGVDGRLQTRGDPPFGQDQNHQYPNEKGFRIDDDPFAN
DGQPIDISDDDLPF 164

FIG. 81

ATGCTGGAACCGTATGGGAAACATTGAAAACGGCTTTCTCCCT
 TTATTTATTATA CGGCAATGAGCCGTTTATTACGGAAACGTATGAGC 100
 GATTGGTGAACCGCAGCGCTGGCCCCGAGGAGCAGGAGTGGAACTTGCT
 GTGTACGACTGCGAGGAAACGCCGATCGAGGCAGCGCTTGAGGAGGCCA 200
 GACGGTGCCTTTCGGCGAGCGCGCTGTCATTCTCATCAAGCATCCAT
 ATTTTTTACGTCTGAAAAGAGAAGGAGATCGAACATGATTGGCGAAG 300
 CTGGAGGCGTACTTGAAGGCAGCGCTCGCCGTTTCGATCGTCGTCTTTT
 CGCGCCGTACGAGAAGCTGATGAGCGAAAAAAATTACGAAGCTGCCA 400
 AAGAGCAAAGCGAAGTCGTATGCCGCCGCTGCCGAAGCGGAGCTG
 CGTGCCTGGGTGCGCGCCGATCGAGAGCCAAGGGCGCAAGCAAGCGA 500
 CGAGGCAGATTGATGTCCTGTTGCAGCGGGACGCAGCTTCCGCC
 TGGCGAATGAAATCGATAATTGGCCCTGTTGCCGGATGGCGGAACC
 ATCGAGGCAGCGCCGTTGAGCGGCTTGTGCCCCCACGCCGAAGAAAAA 600
 CGTATTGTGCTTGTGAGCAAGTGGCGAACCGCAGACATTCCAGCAGCGT
 TGCAGACGTTTATGATCTGCTGAAAACAATGAAGAGCCGATCAAAATT 700
 TTGGCGTTGCTGCCGCCATTCCGCTTGCCTTCGCAAGTGAATGGCT
 TGCCTCTTAGGCTACGGACAGGCAGAACATTGCTGCGCGCTCAAGGTGC 800
 ACCCGTCCCGCTCAAGCTCGCTTGCTCAAGCGGCCGCTCGCTGAC
 GGAGAGCTTGTGAGGCAGTCACAGAGCTCGCTGACGCCATTACGAAGT 900
 GAAAAGCGGGCGGTCGATGCCGGTGGCCGTTGAGCTGCTCTGATGC 1000
 GCTGGGGCGCCCGCCGGCGCAAGCGGGCGCACGCCGGCGG

FIG. 82

MLERVWGNIEKRRFSPLYLLYGNEPFLLTETYERLVNAALGPEEREWNL
 VYDCEETPIEAALEEAETVPFFGERRVILIKHPYFFTSEKEKEIEHDLAK 100
 LEAYLKAPSPFSIVVFFAPYEKLDERKKITKLAKEQSEVVIAPLAEL
 RAWVRRIIESQQAQASDEAIDVLLRAGTQLSALANEIDKLALFAGSGGT 200
 IEAAAVERLVRTPEENVFVLVEQVAKRDIPAAALQTFYDLLENNEEPIKI
 LALLAAHFRLLSQVKWLASLGYQQAQIAAALKVHPRVKLALAQAARFAD 300
 GELAEAINELADADYEVKSGAVDRRLAVELLMRWGARPAQAGRGR

FIG. 83

ATGCGATGGAACAGCTAGC	AAACGCCAGCCGGTGGCGAAAATGCT	
GCAAAGCGGCTTGAAAAGGGCGGATTCTCATGCGTACTTGTGAGG		100
GGCAGCGGGGACGGCAAAAAGCGGCCAGTTGTTGGCGAAACGT		
TTGTTTGTCTGCCCCAATCGGAGTTCCCGTGTAGAGTGCCGCAA		200
CTGCCGGCGCATCGACTCCGGCAACCACCCGTACGTCGGGTGATCGGCC		
CAGATGGAGGATCAATCAAAAGGAACAAATCGAATGGCTGCAGCAAGAG		300
TTCTCGAAAACAGCGGTGAGTCGGATAAAAATGTACATCGTTGAGCA		
CGCCGATCAAATGACGACAAGCGCTGCCAACAGCCTCTGAAATTGG		400
AAGAGCCGCATCCGGGACGGTGGCGTATTGCTGACTGAGCAATACAC		
CGCCTGCTAGGGACGATCGTTCCCGCTGTCAGTGCTTTCGTTCCGGCC		500
GTTGCCGCCGGCAGAGCTCGCCCAGGGACTTGTGAGGAGCACGTGCCGT		
TGCCGTTGGCGCTGTTGGCTGCCATTGACAAACAGCTTCGAGGAAGCA		600
CTGGCGCTTGCCAAGATAGTTGGTTGCCGAGGCGCAACATTAGTGCT		
ACAATGGTATGAGATGCTGGCAAGCCGGAGCTGCAGCTTGTGTTCA		700
TCCACGACCGCTTGTGTTCCGATTGGAAAGCCATCAGCTTGACCTT		
GGACTTG		757

FIG. 84

MRWEQLAKRQPVVAKMLQSGLEKGRISHAYLFEGQRGTGKKAASLLLAKR		
LFCLSPIGVSPCLECRNCRRIDSGNHPDVRVIGPDGGSIKKEQIEWLQQE		100
FSKTAVESDKKMYIVEHADQMTTSAAANSLLKFLEEPHPGTAVAVLTLTEQYH		
RLLGTIVSRCQVLSFRPLPPAELAQGLVEEHVPLPLALLAAHLTNSFEEA		200
LALAKDSWFAEARTLVLQWYEMLGKPELQLLFFIHDRLFPHFLESHQLDL		
GL		252

FIG. 85

GTGGCATACCAAGCGTTATATCGCGTGTTCGGCCGCAGCGCTTGCGGA	100
CATGGTCGGCCAAGAACACGTGACCAAGACGTTGAAAGCGCCCTGCTTC	
AACATAAAAATATCGCACCGCTTAATTATTTCCGGCCCGCGCGGTACAGGA	200
AAAACGAGCGCAGCGAAAATTTCGCCAAGGC GGTC AACTGTGAACAGGC	
GCCAGCGGCGGAGCCATGCAATGAGTGTCCAGCTTGCCTCGGCATTACGA	300
ATGGAACGGTTCCCGATGTGCTGGAAATTGACGCTGCTTCCAACAACCGC	
GTCGATGAAATT CGT GAT ATCCGTGAGAAGGTGAAATTGCGCCAACGTC	400
GGCCCGCTACAAAGTGTATATCATCGACGAGGTGCATATGCTGTCATCG	
GTGCGTTAACCGCGCTGTTGAAAACGTTGGAGGAGCCGCCAAACACGTC	500
ATTTCACTTTGGCCACCGACCGAGCCGCACAAAATTCCGGCGACGATCAT	
TTCCCGCTGCCAACGGTTCGATTTCGCCGCATCCCCTTCAGGCATCG	600
TTTCACGGCTAAAGTACGTCGCAAGCGCCCAAGGTGTCGAGGC GT CAGAT	
GAGGCATTGTCGCCATGCCCGTGC TG CAGACGGGGGATGCGCGATGC	700
GCTCAGCTTGCTTGATCAAGCCATT CGTT CAGCGAACGGAAACTTCGGC	
TCGACGACGTGCTGGCGATGACCGGGGCTGCATCATTTGCCGCTTATCG	800
AGCTTCA TCGAAGCCATCCACCGCAAAGATA CAGCGGCGGTCTTCAGCA	
CTTGGAAACGATGATGGCGCAAGGGAAAGATCCGCATCGTTGGTTGAAG	900
ACTTGATTTGTA CTATCGCATT ATTGCTGTACAAAACCGCTCCCTAT	
GTGGAGGGAGCGATTCAAATTGCTGTCGTTGACGAAGCGTTCACTTCAGT	1000
GTCCGAAAATGATTCCGGTTCCAATT TATA CGAGGCCATCGAGTTGCTGA	
ACAAAAGCCAGCAAGAGATGAAGTGGACAAACCACCCGCCCTCTGTTG	1100
GAAGTGGCGCTTGTGAAACTTTGCCATCCATCAGCCGCCGCCCGTCGCT	
GTCGGCTTCCGAGTTGGAACCGTTGATAAAAGCGGATTGAAACGCTGGAGG	1200
CGGAATTGCGGCCCTGAAGGAACAACCGCCTGCCCTCCGTCGACCGCC	
GCGCCGGTAAAAAAACTGTCCAACCGATGAAAACGGGGGATATAAACG	1300
CCCGGTTGGCCGCATTTACGAGCTGTTGAAACAGGGGACGCATGAAGATT	
TAGCTTGGTGAAGGATGCTGGCGGATGTGCTCGACACGTTGAAACGG	1400
CAGCATAAAGTGTGACCGCTGCCCTGCTGCAAGAGAGCGAGCCGGTTGC	
AGCGAGCGCCTCAGCGTTGTATTAAAATTCAAATACGAAATCCACTGCA	1500
AAATGGCGACCGATCCACAAGTTCGGTCAAAGAAAACGTCGAAGCGATT	
TTGTTGAGCTGACAAACCGCCGCTTGAAATGGTAGCCATTCCGGAGGG	1600
AGAATGGGGAAAAATAAGAGAAGAGTTCATCCGCAATAAGGACGCCATGG	
TGGAAAAAAGCGAAGAAGATCCGTTAATGCCGAAGCGAAGCGGCTGTTT	
GGCGAAGAGCTGATCGAAATTAAAGAA	1677

FIG. 86

VAYQALYRVFRPQRFADMVGQEHTKTLQSALLQHKISHAYLFSGPRGTG	
KTSAAKIFAKAVNCEQAPAAEPCNECPACLGITNGTVPDVLEIDAASNNR	100
VDEIRDIREKVKFAPTSARYKVI IDEVHMLSIGAFNALLKTLEEPKHV	
IFILATTEPHKIPATIISRCQRDFRRIPQAIVSRLKYVASAQGVEASD	200
EALSAIARAADGGMRDALSLLDQAI FSDGKLRLDDVLAMTGAASFAALS	
SFIEAIHRKDAAVLQHLETMMAQGKDPHRLVEDLILYYRDLLLYKTAPY	300
VEGAIQIAVVDEAFTSLSEMI PVSNLYEAIELLNKSQQEMKWTNHPRLLL	
EVALVKLCHPSAAAPSLSASELEPLIKRIETLEAEIERRLKEOPPAPPSTA	400
APVKKLSKPMKTGGYKAPVGRIYELLKQATHEDLALVKGCADVLDTLKR	
QHKVSHAALLQESEPVAASASAFLVKFKYEIHCKMATDPTSSVKENVEAI	500
LFELTNRRFEMVAIPEGEWGKIREEFIRNKDAMVEKSEEDPLIAEAKRLF	
GEELIEIKE	559

FIG. 87

ATGGTGACAAAAGAGCAAAAGAGCGGTTCTCATCCTGTTGAGCAGCT	
GAAGATGACGTGGACGAATGGATGCCGATTTCGTGAGGCAGCCATTC	100
GCAAAGTCGTATCGATAAAAGAGGAGAAAAGCTGGCATTTTATTTCA	
TTCGACAACGTGCTGCCGGTCATGTATACAAAACGTTGCCATCGGCT	200
GCAGACGGCGTTCCGCCATATCGCCGCCGTCGCCATACGATGGAGGTG	
AAGCGCCGCGCTAAGTGAAGGCGATGTGCAGGCGTATTGGCCGTTGC	300
CTTGGCGAGCTGCAAGAAGGCATGTCGCCGTTGTCAGTGGCTCAGCCG	
GCAGACGCGCTGAGCTGAAAGGAAACAAGCTGCTTGTGCGTTGCCGCC	400
AAGCGGAAGCGCTGGCGATCAAACGGCGGTCGCCAAAAAAATCGCTGAT	
GTGTACGCTTCGTTGGGTTCCCCCTTCAGCTGACGTAGCGTCGA	500
GCCGTCCAAGCAAGAAATGGAACAGTTTGGCGCAAAAACAGCAAGAGG	
ACGAAGAGCGAGCGCTTGCCTACTGACCGATTAGCGAGGGAAAGAAGAA	600
AAGGCCGCGTCTGCCGCCGTCGGTCCGCTTGTATCGGCTATCCGAT	
CCGCGACGAGGAGCCGGTGCGGCGGCTGAAACGATCGTGAAGAAGAGC	700
GGCGCGTCGTTGTCAAGGCTATGTATTGACGCCAGTGAGCGAATT	
AAAAGCGGCCGACGCTGTTGACCATGAAAATCACAGATTACACGAAC	800
GATTTAGTCAAAATGTTCTCGCGCAGAAAGAGGACGCCGAGCTTATG	
GGCGCGTCAAAAAGGCATGTGGGTGAAAGTGCAGCGCAGCGTGC	900
GATACGTTCGTCGTGATTTGGTCATCATGCCAACGATTGAAACGAAAT	
CGCCGCAAACGAACGGCAAGATAACGGCGCCGAAGGGAAAAGAGGGT	1000
AGCTCCATTGCAACCCCCGATGAGCCAAATGGACGCCGTCACCTCGGT	
ACAAAACCTATTGAGCAAGCGAAAAAAATGGGGGCATCCGGCGATCGCC	1100
CACCGACCATGCCGTTGTCAGCTGTTCCGGAGGCCTACAGCGCGGCA	
AAAAACACGGCATGAAGGTCAATTACGGCCTTGAGGCCAACATCGTC	1200
GATGGCGTGCCGATGCCCTACAATGAGACGCCGCGCTTTCGGAGGA	
AACGTACGTCGTCTTGACGTCGAGACGCCGCGCTGCGGCTGTGTACA	1300
ATACGATCATTGAGCTGGCGCGGTGAAAGTGAAGACGGCGAGATCATC	
GACCGATTGATGCGTTGCCAACCTGGACATCCGTTGCGGTGACAAC	1400
GATGGAGCTGACTGGGATACCGATGAGATGGTGAAGACGCCCGAAC	
CGGACGAGGTGCTAGCCGTTTGTGACTGGGCCGGCATGCGACGCTT	1500
GTTGCCACACGCCAGCTTGACATCGGTTAAACGCCGGCCTCGC	
TCGCATGGGGCGCGCAAATCGCAATCCAGTCATCGATACGCTCGAGC	1600
TGGCCCCTTTTATACCCGGATTGAAAAACCATCGCTCAATACATTG	
TGCAAAAATTTGACATTGAATTGACGCACTACCGCGCCATCTACGA	1700
CGCGGAGGCGACCGGGCATTGCTTATGCCGCTGTTGAAGGAAGCGGAAG	
AGCGCGCATACTGTTCATGACGAATTAAACGCCGCACGCACAGCGAA	1800
GCGTCCTATCGCTTGCGGCCGTTCCATGTGACGCTGTTGGCGCAAA	
CGAGACTGGATTGAAAATTGTTCAAGCTGTGTCATTGTCACATTC	1900
AATATTTCACCGTGTGCCGCGCATCCCGCGCTCCGTGCTCGTCAAGCAC	
CGCGACGGCCTGCTGTCGGCTCGGGCTCGACAAAGGAGAGCTGTTGA	2000
CAACTTGATCCAAAAGGCCGGAGAAGTGAAGACATGCCCGTTTT	
ACGATTTCCTGAAAGTGCATCCGCCGGACGTGTACAAGGCCGCTCATCG	2100
ATGGATTATGTGAAAGACGAAGAGATGATCAAAACATCATCGCAGCAT	
CGTCGCCCTGGTGAGAAGCTTGACATCCGGTTGTCGCCACTGGCAACG	2200

FIG. 88A

TCCATTACTGAACCCAGAAGATAAAATTACCGGAAAATCTTAATCCAT	
TCGCAAGGCAGGGCGAATCCGCTCAACCGCCATGAACGTGCCGATGTATA	2300
TTTCCGTACGACGAATGAAATGCTGACTGCTCTCGTTTAGGGCCGG	
AAAAAGCGAAGGAAATCGTCGTGACAACACGCAAAAATCGCTCGTTA	2400
ATCGGCAGATGTCAAGCCGATCAAAGATGAGCTGTATACGCCGCGCATTGA	
AGGGGCGGACGAGGAAATCAGGGAAATGAGCTACCGCGGGCGAAGGAAA	2500
TTTACGGCAGCCGTTGCCGAAACTTGTGAAGAGCGGCTTGAGAAGGAG	
CTAAAAAGCATCATCGGCCATGGCTTGCCGTCAATTATTTGATCTCGCA	2600
CAAGCTTGTGAAAAAAATCGCTCGATGACGGCTACCTGTGCGGTCGCG	
GATCGGTCGGCTCGTGTGCGACGATGACGGAATCACCGAGGTC	2700
AATCCGCTGCCGCCGATTACGTTGCCGAAC TGCAAGCATTGGAGTT	
CTTTAACGACGGTTCACTCGGCTCAGGGTTGATTGCCGATAAAAAC	2800
GCCCAGATGTGGGACGAAATACAAGAAAGACGGGACGACATCCCCTT	
GAGACGTTTCTCGGTTAAAGGCGACAAAGTGCCGATATCGACTTGAA	2900
CTTTTCCGGCGAATACCAGCCGCGCCCACAAC TACGAAAGTGTGT	
TTGGCGAAGACAAACGTCTACCGCGCCGGACGATTGGCACGGTCGCTGAC	3000
AAAACGGCGTACGGATTGTCAAAGCGTATGCGAGCGACCATAACTTAGA	
GCTGCGCGCGCGGAAATCGACGGCTCGCGCTGGCTGCACCGGGGTGAA	3100
GCGGACGACCGGGCAGCATCGGGCGCATCGTCGCCCCGATTATA	
TGGAAATTACGATTTCAGCCGATTCAATATCCGGCCGATGACACGTCC	3200
TCTGAATGGCGGACGACCCATT CGACTTCCATT CGATCCACGACAATT	
GTTGAAGCTCGATATTCTCGGCACGACGATCCGACGGTCATT CGCATGC	3300
TGCAAGATTAAAGCGGATCGATCCGAAACGATCCCACCGACGACCCG	
GATGTGATGGCATTTCAGCAGCACCGAGCCGCTGGCGTTACGCCGGA	3400
GCAAATCATGTGCAATGTCGGCACGATCGGATTCCGGAGTTGGCACGC	
GCTTCGTTGGCAAATGTTGGAAGAGACAAAGGCCAAAACGTTTCCGAA	3500
CTCGTGC AATTTCGGCTGTGCGCACGGCACCGATGTGTGGCTCGGCAA	
CGCGCAAGAGCTCATTCAAACGGCACGTGTACGTTATCGGAAGTCATCG	3600
GCTGCCGCGACGACATTATGGTCTATTGATTACCGCGGCTCGAGCCG	
TCGCTCGCTTTAAAATCATGGAATCCGTGCGCAAAGGAAAAGGCTTAAC	3700
GCCGGAGTTGAAAGCAGAAATGCGCAAACATGACGTGCCGGAGTGGTACA	
TCGATT CATGCAAAAAATCAAGTACATGTTCCGAAAGCGCACGCCGCC	3800
GCCTACGTGTTAATGGCGGTGCGCATCGCCTACTTTAAGGTGCACCATCC	
GCTTTGTATTACCGCTCGTACTTACGGTGCAGGGCGGAGGACTTGACC	3900
TTGACGCCATGATCAAAGGATCACCCGCCATT CGCAAGCGGATTGAGGAA	
ATCAACGCCAAAGGCATT CAGGCAGGGCGAAAGAAAAAGCTTGCTCAC	4000
GGTTCTTGAGGTGGCTTAGAGATGTGCGAGCGCGGCTTCCCTTAAAA	
ATATCGATTGTACCGCTCGCAGGCAGGGATT CGTCACTGACGGCAAT	4100
TCTCTCATTCCGCCGTTCAACGCCATTCCGGGGCTTGGGACGAACGTGGC	
GCAGGGCGATCGTGCAGGGCGAGGAAGGGAGTTTTGTCGAAGGAGG	4200
ATTTGCAACAGCGCGGCAAATTGTCGAAAACGCTGCTCGAGTATCTAGAA	
AGCCGCGGCTGCCCTGACTCGCTCCAGACCATACCAGCTGCGCTGTT	4300

T

FIG. 88B

MVTKEQKERFLILLEQLKMTSDEWMPHFREAAIRKVVIDKEEKSWHFYFQ 100
 FDNVLPVHVYKTfadRLQTAFRHIAAVRHTMEVEAPRVTEADVQAYWPLC
 LAELQEGLMSPLVDWLRSQTPELKGKLLVVARHEAEALAIKRRFAKKIAD
 VYASFGFPPLQLDVSEPSKQEMEQFLAQKQQEDEERALAVLTDLAREEE 200
 KAASAPPSPGPLVIGYPIRDEEPVRRLETIVEERRVVQGYVFDAEVSEL
 KSGRTLLTMKITDYTNSILVKMFSRDKEDAEMLMSGVKKGWMVKVRGSVQN 300
 DTFVRDLVIIANDNEIAANERQDTAPEGEKRVELLHTPMSQMDAVTSV
 TKLIEQAKKWGHPAIAVTDHAVVQSFPEAYSAAKKHGMKVIYGLEANIVD
 DGVPIAYNETHRRLSEETYVVFVDTETGLSAVYNTIELAAVKVKDGEII 400
 DRFMSFANPGHPLSVTTMELTGITDEMVKDAPKDEVLARFVDWAGDATL
 VAHNASFIDGFLNAGLARMGRGKIANPVIDTLELARFLYPDLKNHRLNTL
 CKKFDIELTQHHRAIYDAEATGHILLMRLLKEAEERGILFHDELNSRTHSE 500
 ASYRLARPFTVTLAQNGETGLKNLFKLVSLSHIQYHRVPRIPRSVLVKH
 RDGLLVSGSGCDKGELFDNLIQKAPEEVEDIARFYDFLEVHPPDVYKPLIE 600
 MDYVKDEEMIKNIIRSIVALGEKLDIPVVAATGNVHYLNPEDKIYRKILIH
 SQGGANPLNRHELPDVYFRTTNEMLDCFSFLGPEKAKEIYGDPLPKLVEERLEKE 700
 IGDVKPIKDELYTPRIEGADEEI REMSYRRAKEIYGDPLPKLVEERLEKE
 LKSIIGHGFAVIYLISHKLVKKSLDDGVLVGSRGSGVGSSFVATMTEITEV 800
 NPLPPHYVCNPCKHSEFFNDGSVGSGFDLDPDKNCPRCGTKYKKDGHDIPE 900
 ETFLGFKGDKVPDIDLNFSGEYQPRAHNYTKVLFGEDNVYRAGTIGTVAD 1000
 KTAYGFVKAYASDHNLLELRGAEIDLAACTGVKRTTGQHPGGIIVVPDYM
 EIYDFTPPIQYPADDSSEWRTHFDFHSIHNDNLLKLDILGHDDPTVIRML 1100
 QDLSGIDPKTIPTDDPDVMGIFSSTEPLGVTPEQIMCNVGTIGIPEFGTR
 FVRQMЛЕЕTRPKTFSELVQISGLSHGTDVWLGNQAQELIQNGTCTLSEVIG 1200
 CRDDIMVYLIYRGLEPSLAFKIMESVRKGKGLTPFEAEMRKHDVPEWYI
 DSCKKIKYMFPAKAHAAAYVLMAVRIAYFKVHHPLLYYASYFTVRAEDFDL 1300
 DAMIKGSPAIRKRIEEINAKGIQATAKEKSLLTVLEVALEM CERGFSFKN
 IDLYRSQATEFVIDGNSLIPPFNAIPGLGTNVQAIVRAREEEGEFLSKED 1400
 LQQRGKLSKTLLEYLESRGCLDSL PDHNQLSLF

FIG. 89